

Toxicological stress indicators in human red blood cells : changes in glutathione and glutathione S-transferase as biological markers for electrophilic and oxidative stress

Citation for published version (APA):

Evelo, C. T. A. (1994). *Toxicological stress indicators in human red blood cells : changes in glutathione and glutathione S-transferase as biological markers for electrophilic and oxidative stress*. [Doctoral Thesis, Maastricht University]. Rijksuniversiteit Limburg. <https://doi.org/10.26481/dis.19950119ce>

Document status and date:

Published: 01/01/1994

DOI:

[10.26481/dis.19950119ce](https://doi.org/10.26481/dis.19950119ce)

Document Version:

Publisher's PDF, also known as Version of record

Please check the document version of this publication:

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Toxicological Stress Indicators in Human Red Blood Cells

**Changes in Glutathione and Glutathione S-Transferase
as Biological Markers
for Electrophilic and Oxidative Stress**

CIP-DATA Koninklijke Bibliotheek, Den Haag

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Toxicological stress indicators in human red blood cells :
Changes in glutathione and glutathione S-transferase as
biological markers for electrophilic and oxidative stress. /
Christoffel Theodorus Anthonius Evelo. – Maastricht :
Universitaire Pers Maastricht. – Ill.

Thesis Maastricht. – With ref. – With summary in Dutch.

ISBN 90-5278-168-0

Subject headings: toxicology / erythrocytes / glutathione.

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Financial support by the corporate safety, health, environment and technology division of DSM, the Dr.Ir. J.H.J. van de Laar foundation for the advancement of biochemical research and by Perkin-Elmer Nederland B.V. and Bio-Rad Laboratories B.V. for the publication of this thesis is gratefully acknowledged.

Toxicological Stress Indicators in Human Red Blood Cells

**Changes in Glutathione and Glutathione S-Transferase
as Biological Markers
for Electrophilic and Oxidative Stress**

PROEFSCHRIFT

ter verkrijging van de graad van doctor
aan de Rijksuniversiteit Limburg te Maastricht,
op gezag van de Rector Magnificus, Prof.Dr. H. Philipsen,
volgens het besluit van het College van Dekanen,
in het openbaar te verdedigen
op donderdag 19 januari 1995 om 16.00 uur

door

Christoffel Theodorus Anthonius Evelo

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The investigations presented in this thesis were carried out at the department of Toxicology of the university of Nijmegen (chapters 3 and 7), at the former department of Occupational & Environmental Health and Toxicology (chapters 6, 8 and 9) and at the department of Pharmacology (Toxicology section) of the university of Limburg (chapters 4 and 5).

“It is quite imaginable that scientists abandon a paradigm out of frustration and not because they have arguments against it.

...

Besides, it is not at all prudent to put too much trust in experimental results. Indeed, it would be a complete surprise and even a cause for suspicion, if all the available evidence should turn out to support a single theory, even if this theory happened to be true”

Paul Feyerabend

in: Criticism and the growth of knowledge

Voor Chantal, Jannemiek en Sjoerd

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Abbreviations used/Gebruikte afkortingen:

3-HAA	3-hydroxyacetanilide
4-PDS	4,4'-dithiodipyridine
5-FU	5-fluorouracil
6PGDH	6-phospho-gluconate dehydrogenase
β -Gal	β -galactosidase
β_2 -M-S	β_2 -microglobulin in serum
β_2 -M-U	β_2 -microglobulin in urine
AAP	alanine aminopeptidase
ALAT	alanine aminotransferase
ALB-U	albumine in urine
ALP	alkaline phosphatase
AOS	active oxygen species
Ara-C	cytosine arabinoside
ASAT	aspartate aminotransferase
Bil-tot	total bilirubin
BM	biological monitoring
BEM	biological effect monitoring
c-DDP	cisplatin (cis-diamminedichloroplatinum(II))
CBB	Coomassie Brilliant Blue
CDNB	1-chloro-2,4-dinitrobenzene
CP	cyclophosphamide
Creat-S	creatinine in serum
DCP	1,3-dichloropropene
DEM	diethyl maleate
DTT	dithiothreitol
DTNB	5,5'-dithio-bis(2-nitrobenzoic acid)
EDTA	ethylenediaminetetraacetic acid
ELISA	enzyme linked immunosorbent assay
EM	environmental monitoring
F6P	fructose 6-phosphate
FAD	flavin adenine dinucleotide
G6P	glucose 6-phosphate
G6PDH	glucose 6-phosphate dehydrogenase
GPX	glutathione peroxidase
GSH	glutathione (reduced)
GSSG	glutathione (oxidized)
GST	glutathione S-transferase
GR	glutathione reductase
GR _{coeff}	GR riboflavin activity coefficient
GGT	γ -glutamyltranspeptidase
GT	glutathione (total)

Hb	hemoglobin
Hb ⁺	methemoglobin: hemoglobin with iron in the ferric state
HbSH	free sulfhydryl groups in hemoglobin
HS	health surveillance
HYAM	hydroxylamine (HONH ₂)
IAcA	iodoacetamide
IEF	isoelectric focusing
FPLC	fast protein liquid chromatography
LP	lipid peroxidation
MAS	microsomal activating system
MDA	malondialdehyde
MTX	methotrexate
NAD(H)	β -nicotinamide adenine dinucleotide (H=reduced)
NADH-HbR	NADH-methemoglobin reductase
NADP(H)	β -nicotinamide adenine dinucleotide phosphate (H=reduced)
NADPH-HbR	NADPH-methemoglobin reductase
NEM	N-ethylmaleimide
NODMH	N,O-dimethyl hydroxylamine (CH ₃ ONHCH ₃)
OEH	O-ethyl hydroxylamine (CH ₃ CH ₂ ONH ₂)
OEL	occupational exposure limit (Dutch)
OMH	O-methyl hydroxylamine
PBS	phosphate-buffered saline
RBC	red blood cells
RBP	retinol binding protein
RS	NADPH regenerating system
SD	standard deviation
SE	standard error
SEMD	standard error of the mean difference
SLDH	serum lactate dehydrogenase
TBA	thiobarbituric-acid
TBARS	thiobarbituric-acid reactive substance
TCA	trichloroacetic acid
TLV	threshold limit value
TT	thioltransferase

Chapter 1

Outline

This thesis describes the use of changes in glutathione (*GSH* or *GT*)¹ levels in human blood and glutathione S-transferase (*GST*) activity in human red blood cells for the assessment of electrophilic and oxidative stress. For the study of biological effects of chemical exposure in man, red blood cells form a rather unique entity. For routine purposes blood cells are the only cell type from within the body that can be made easily available, and red blood cells are by far the most abundant blood cells. Red cells are also very special cells. They do not contain a nucleus or mitochondria. This, combined with the very high oxygen turnover and special membrane properties, leads to a special biochemistry. As it will be discussed in chapter 2, both *GSH* and *GST* play a prominent part in erythrocyte biochemistry. This prominence combined with the highly specialized physiology of the erythrocyte may however result in increased vulnerability of both *GSH* and *GST*, which may in turn be of use for biological effect monitoring.

In chapter 2 a general introduction to the subject is given. The biochemistry of red blood cells and the physiological functionality of *GSH* and *GSTs* are discussed. Chapters 3 to 5 are descriptions of *in vitro* and *in vivo* laboratory experiments to study the vulnerability of *GSH* and *GSTs* during electrophilic and oxidative stress. In chapter 3 depletion of *GT* after *in vitro* treatment of human red blood cells with some model electrophilic compounds is compared with another well-known effect parameter, the formation of hemoglobin adducts. In chapter 4 the *in vitro* effects of a number of cytostatic drugs on *GT* and *GST* in erythrocytes are described. Since the efficacy of some of these drugs depends on metabolic activation by membrane bound mixed function oxygenases the *in vitro* system was extended with a rat derived liver microsomal activating system.

1. Throughout this thesis the abbreviations *GSH*, *GSSG* and *GT* are used to discriminate reduced, oxidized and total glutathione respectively. The abbreviation *GSSG* is only used to explicitly indicate oxidized glutathione, as is usual in biochemical literature. The abbreviation *GT* is less usual, but also uncomplicated. *GT* explicitly indicates the total of reduced and oxidized glutathione, as it is determined with the glutathione reductase dependent cyclic oxidation-reduction assay. The use of *GSH* in the literature, and in some cases in this thesis, is unfortunately rather ambivalent. It is both used to indicate reduced glutathione (i.e. with a free sulfhydryl (SH) group) and as a general abbreviation for glutathione.

In chapter 5 an extensive *in vitro* study towards the hematotoxicity of three industrially used hydroxylamines is described. Next to effects on glutathione and GST, several other biochemical effects were studied.

In the next part — chapter 6–9 — several human monitoring studies are described. Chapter 6 is supplied as an introduction to the use of biological effect monitoring methods in occupational toxicology. It also contains some descriptions of pilot studies in smokers and creosote workers, and reviews part of the results described in the following chapters. Chapter 7 describes a biological effect monitoring study in pesticide applicators using the soil fumigant 1,3-dichloropropene. In these workers classical liver and kidney toxicity tests were also performed. This allows for an interesting comparison of the sensitivity of the methods used. In chapter 8 a study towards the effects of long distance running is described. Physical activity leads to increased oxygen turnover rates and to increases in the formation of active oxygen species. This chapter describes that the GT and GST parameters can be used to assess the increased oxidative stress resulting from contest participation. The changes in basal values of these and some related parameters during endurance training were also studied. Finally, chapter 9 describes the effects found in ex-miners with coal workers pneumociosis. Here oxidative stress probably results from the continuous inflammatory activity in the lung.

The findings from chapter 3 to 9 are summarized and discussed in chapter 10. A summary in Dutch is included in chapter 11 (*Een samenvatting in het Nederlands is te vinden in hoofdstuk 11*).

Chapters 3, 6, 7, 8 and 9 were published previously (see the respective title pages for details). In some cases minor modifications such as the completion of references in press at the time of publication, correction of typing errors and addition of newer relevant references were made.

Chapter 2

General Introduction

As occupational toxicologists we face an intriguing dilemma. When healthy persons are exposed to low quantities of noxious chemicals small changes in some biological parameters may occur. To know the nature and magnitude of such changes could be very helpful in estimations of the exposure that has occurred and in evaluations of a possible resulting risk [65, 168]. The problem is that interesting changes often occur in tissues that are not accessible for routine analyses. Pesticide applicators, nurses handling cytostatic drugs, process operators in chemical industry and many others exposed to chemicals in their daily work may all welcome our interest in their personal welfare, and show a positive interest in our work. Their enthusiasm would definitely transform into a more defensive attitude when we would ask them for tissue samples. The use of for instance liver tissue may be useful in animal studies and even in some *in vitro* experimental studies using human tissue; it is not a practical option in occupational toxicology. What remains for us is mainly the determination of chemicals or their metabolites in excreta and in some cases in clippings like nails and hair. The problem is that such so called “biological monitoring” studies [17] do give us an indication of the exposure that has occurred, but not of any possible effect that resulted of it. As a matter of fact this problem is not limited to toxicology. Clinicians have always had an interest in possible changes in the functionality of organs buried deep in the human body. And, more often than not, they too encounter situations where cutting is not the most appropriate way of doing. Fortunately, alternatives were found. In many cases disturbances of internal physiological functions are visible from the outside or can be monitored using readily available samples of excreta and blood. Well-known examples of the use of urine and blood samples are the determination of liver and kidney function, of genetic variations, of hematological disorders, of factors of immunological importance and of aberrations of hormonal regulation including diabetes. In fact the possibilities to monitor health state using biological samples are so extensive that they are covered by a separate discipline: clinical chemistry. Books describing the multitude of tools available to the clinical chemist (e.g. [71, 136, 154]) are also a most valuable source of information for the occupational toxicologist [55, 84, 168].

An important difference of interest between clinicians and occupational toxicologists should however be noted. The clinician is primarily interested in parameters that may indicate changes in health state. Monitoring of health effects (or health surveillance [168]) is of course also of interest to the toxicologist, although it belongs more to the domain of the occupational physician. As stated, the toxicologist is primarily interested in biomarkers¹ that might indicate the occurrence of exposure to hazardous compounds or that might be useful for evaluation of the risks resulting thereof. For both purposes it is not necessary that the parameters evaluated are in itself a consequence of impaired health. For this reason we may use clinical parameters at a level where they are not of clinical significance, or choose to use entirely different parameters.

Next to conceptual difficulties [134,168], the use of clinical parameters for the study of early effects introduces some problems of methodological nature. The actual changes in the parameters determined are often very small, and the values affected may still lay within the normal range. This means that often sensible studies can only be done at a group level, which is in contrast with the more individual centered approach of the physician. Another useful approach is not to study just the momentary value of a parameter but to look at its change in time. Most often this is done by comparison of pre- and post-exposure values. In this way much more sensitive methods can be developed, assisted at large by the ability to use paired comparison statistics. On the other hand, the use of comparisons of multiple analyses performed at different times, asks for rigid control of between assay variations and seasonal changes [55]. In general such controls can be done by the inclusion of control samples with known values in each analyzed batch of samples. Again the final analyses can be performed on a group basis (i.e. statistical analyses to verify whether *changes* in values are significant), or on an individual basis. In the latter case differences between toxicological and clinical analyses will tend to disappear again, as many clinicians will argue that significant individual changes within the normal range might still be of clinical value.

Since biomarkers used in toxicology need not always be indicative of individual health changes, parameters that are not useful to the clinician may prove to be of use for the toxicologist. Especially when monitoring is performed to evaluate the risk of quantal effects like the development of cancer, the use of such early markers is indicated. Good examples of this approach can be seen in the determination of nucleic acid [117,161] or protein adducts [46,105] for the monitoring of exposure to electrophilic genotoxic compounds [2,38,163] and in the monitoring of lipid peroxidation products [52,77,97,111,116] as a measure for oxidative stress [12,13,35,49,76,144,156]

1. The term biomarkers—a contraction of *biological markers*—is used here to indicate both substance related parameters—as used in biological monitoring—and effect related parameters—as used in biological effect monitoring and health surveillance—. See chapter 6 for definitions of these three monitoring concepts.

Erythrocytes

Blood cells are of special interest for the assessment of *in vivo* changes in biological parameters. In general, blood cells are the easiest available cells from within the body. They can be obtained with hardly invasive techniques, and therefore volunteers can easily be found. Like it is generally the case for use of human material in biomedical experiments, it should be noted that microbiological contaminations can be present, and special care should be taken with regard to possible viral infections (especially HIV and hepatitis). Human erythrocytes have a longevity of about 120 days [30], and new cells are continuously being formed in the hematopoietic tissues located mainly in the bone marrow. Erythrocytes are extremely specialized cells. Their main function is the transport of oxygen from the lungs to peripheral tissues. To enable them for this task a very large fraction (about 80% of the soluble protein) of the erythrocyte consists of hemoglobin. Other adaptations result from the necessity to be able to squeeze through tiny spaces between cells. For this reason erythrocyte membranes are very flexible. This is attained by a high abundance of poly-unsaturated fatty acids, and by specialized membrane proteins. Also, many organelles normally present in human cells are removed from the erythrocyte after the process of erythropoiesis, and mature erythrocytes do no longer contain a nucleus, mitochondria or ribosomes. As a consequence erythrocytes do not contain any DNA and are not able to divide or to synthesize new proteins. In effect erythrocytes are slowly dying cells [30]. Because of the absence of mitochondria the erythrocyte is fully dependent on glycolysis for phosphorylation of ATP and reduction of NAD^+ to NADH and on the so called hexose monophosphate shunt for reduction of NADP to NADPH. The key enzyme in this hexose monophosphate shunt is glucose 6-phosphate dehydrogenase.

The absence of *de novo* protein synthesis in mature erythrocytes has important consequences for the usability of erythrocytes for monitoring purposes. In most other cell types damaged proteins will be removed and resynthesized. While erythrocytes do contain proteolytic enzymes which are able to remove oxidized proteins [36], hemoglobin adducts are not normally removed [46]. For the monitoring of hemoglobin adducts this is a great advantage, especially when compared with the monitoring of DNA adducts. While DNA adducts are rapidly repaired, the hemoglobin adducts remain present in the erythrocyte until sequestration of this cell. Therefore, it is possible to use hemoglobin adduct measurements some time after incidental exposure has occurred—in principle up to 3 months—or to determine a cumulative dose as a result of chronic exposure. Sequestration of erythrocytes is not a simple age related phenomenon. A fraction of the erythrocytes is removed independent of age. In rats this random destruction appears to be of more importance than in man [105], and amounts to about 3% of the erythrocytes per day. In such cases the removal of erythrocytes can best be described by a function that combines exponential removal of old cells with random

removal [72]. When changes in activities of enzymes present in erythrocytes are used as an effect parameter, the stability of the affected enzymes is not even a prerequisite. Since irreversible inactivation of erythrocyte enzymes will lead to decreased activities, older erythrocytes are expected to have lower activities of vulnerable enzymes. In accordance with this expectation, an age related decline in activities of aspartate amino transferase [114], glutathione reductase [73,114,148], glutathione peroxidase [73], glucose 6-phosphate dehydrogenase [73], and superoxide dismutase [30] has been found in human erythrocytes, while the activity of the erythrocyte glutathione S-transferase (*GST*) studied in this thesis is also known to be affected [50]. It is important not to confuse this decreased activity changes in older cells with the changes occurring in older individuals. Lowered activities of copper-zinc superoxide dismutase, glutathione reductase and *GST* were found in erythrocytes in elderly [26], the same study showed that glutathione peroxidase activity is increased in older persons.

Glutathione

The tripeptide glutathione (*GSH*, L- γ -glutamyl-L-cysteinyl-glycine, see figure 2.1 on the facing page) plays a central role in the protection of cells against oxidative² [75,129,135] and electrophilic stress³ [37,67,135] and radiation [24]. *GSH* is the most abundant non-protein sulfhydryl present in cells, with intracellular concentrations ranging from 0.5 to 10 mM [129].

The synthesis of *GSH* by aerobic cells can be considered as a way to hold high levels of usable cysteine in a non-toxic form. Cysteine itself will rapidly autoxidize in the presence of metal ions like Fe^{3+} and Cu^{2+} [157]. As a result of the hydrogen peroxide liberated during this autoxidation, simple thiols like cysteamine [147,160] and cysteine [160] can be toxic to cells. Blocking of the cysteine amino group by forming γ -glutamyl-cysteine slows its autoxidation considerably [145]. This may have been the selective factor that led to the evolution of γ -glutamylcysteine synthetase [48]. The special γ -binding makes this dipeptide, and glutathione, resistant to normal peptidase activity [44], and γ -glutamyl transpeptidase (γGT ; see figure 2.2) is therefore needed for cleavage of glutathione or glutathione conjugates. Coupling of glycine to the γ -glutamylcysteine dipeptide by glutathione synthetase, leading to *GSH* synthesis, makes it even more resistant to autoxidation [145]. Furthermore, it provides a more extended

2. Oxidative stress refers to the unusually high presence of molecules with a high potency to abstract electrons from biomolecules. In practice free radicals are the most importance class of biological oxidators.

3. Electrophilic stress is defined as the effect of compounds with a high reactivity towards nucleophilic centers like —with increasing reactivity— oxygen, nitrogen and sulphur present in biomolecules. Reaction of electrophiles with nucleophilic centers in biomolecules leads to the formation of new chemicals bonds between the two (adduct formation).

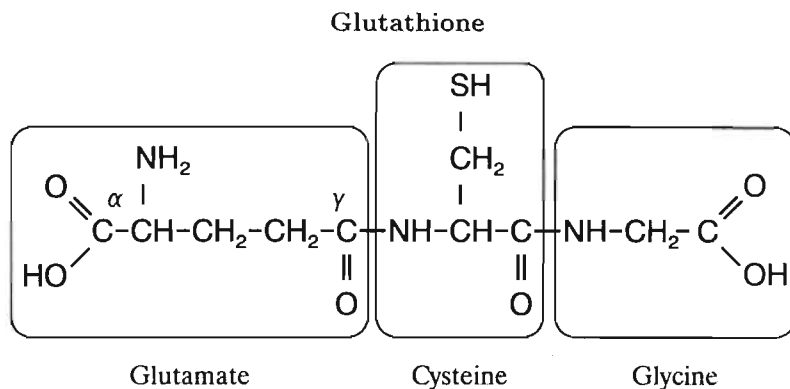


Figure 2.1: The structure of the tripeptide glutathione (GSH: L- γ -glutamyl-L-cysteinyl-glycine). Note the γ linkage between glutamate and cysteine which makes glutathione resistant to normal peptidase activity.

carboxylic site that is used by many glutathione-dependent enzymes for substrate recognition and active site handling [44, 45].

The abundancy of glutathione in biological systems has led to enormous amounts of research to find “the role of glutathione”. The crusade for this Holy Grail—as Kenneth Douglas called it in one of his excellent reviews [44]—led to the recognition of many functions. So many, that the said review was introduced with the quotation:

*There was an old woman who lived in a shoe,
Who had so many children she didn't know what to do.*

Whether the original function of glutathione was to create a save storage form of cysteine or not, currently a myriad of functions is known, and many of them are vital. Some of the more important functions include the detoxification of xenobiotics [91], removal of hydrogen peroxide and other peroxides and free radicals [129], maintenance of free protein sulfhydryl groups [74] and the synthesis of leukotriene LTC₄ and derivatives [70]. These activities may lead to variations in the cellular availability of GSH [159]. This thesis will focus on glutathione loss in erythrocytes as a result of electrophilic or oxidative stress, and on concomitant changes in GST activity.

Due to the high nucleophilic potency of the glutathione sulfhydryl, GSH can function as an excellent nucleophilic scavenger for many reactive electrophiles [37, 45]. In many cases this reactivity is increased further by glutathione S-transferases [91]. Since electrophilic chemicals can also react with biological macromolecules like DNA, and thereby present a genotoxic risk [101], this glutathione conjugation furnishes an important detoxification mechanism. In some

instances however, the glutathione conjugates themselves may be toxic to the cells [5, 39, 103]. Glutathione S-conjugates can be metabolised to the corresponding cysteine conjugates by γ GT and cysteinylglycine dipeptidase and aminopeptidase M [39], which can subsequently be N-acetylated by N-acetyl transferase to yield the mercapturic acid [22, 163] (Figure 2.2). Formation of glutathione conjugates will therefore lead to loss of available glutathione. The thioethers formed are often excreted in urine, and can themselves be used for biological monitoring of exposure to electrophilic compounds [64, 163].

The mercapturic acid pathway.

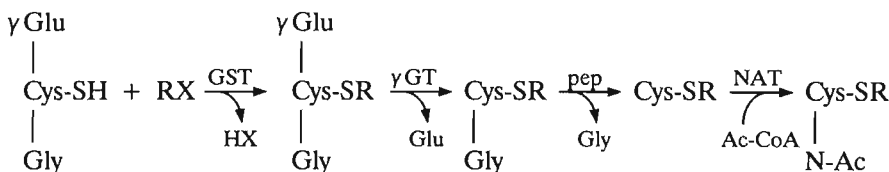
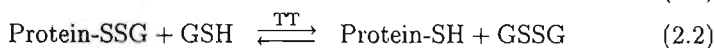
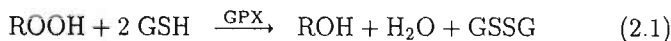


Figure 2.2: Glutathione conjugation by glutathione S-transferase (GST) and subsequent steps of the mercapturic acid pathway. The γ -glutamyl moiety is removed by γ -glutamyl transferase (γ GT). The remaining dipeptide can subsequently be cleaved by cysteinyl glycine dipeptidase or aminopeptidase M to yield a cysteine conjugate. This cysteine conjugate can be N-acetylated by N-acetyl transferase (NAT), which leads to mercapturic acid formation. Note that erythrocytes do not contain γ GT and therefore have to excrete the complete glutathione conjugate.

Upon oxidative stress conditions a variety of processes can occur which may lead to the oxidation of GSH to its oxidized form, the disulfide GSSG. Among those are the activity of glutathione peroxidase [eq. 2.1] and some of the reactions catalysed by thioltransferase [164] [see eq. 2.2 for an example]. A more extensive



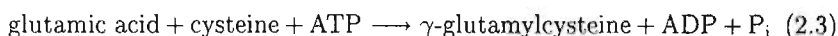
The glutathione peroxidase (GPX) reaction [eq. 2.1] and an example of a thioltransferase (TT) [164] catalysed reaction [eq. 2.2]. Both reactions lead to the oxidation of GSH to GSSG.

description of the role of glutathione in the protection against oxidative stress will be given below. For the moment it suffices to note that such conditions may lead to oxidation of glutathione. In the next section it will be shown that when such an oxidation occurs in erythrocytes the result may be excretion of GSSG.

Glutathione Turnover in Erythrocytes

Oxidized glutathione is exported from erythrocytes by two separate ATP dependent transport systems [19, 81] with apparent K_m values for GSSG of 0.1 mM and 7.1 mM respectively. The two systems are probably related but not identical ATPases with unidirectional transport activity [3, 83]. The high affinity (low K_m) system has a V_{max} of 0.33 nmol/min/ml cells and seems to be devoted to GSSG export. The low affinity system has a ten times higher V_{max} and is also involved in the transport of thioethers formed by conjugation of xenobiotic compounds like 1-chloro-2,4-dinitrobenzene. Awasthi and coworkers [11] demonstrated that thioether export is likely to be the main function of this system, since its GSSG exporting activity is inhibited by thioethers while thioether export is not inhibited by GSSG concentrations up to 11.8 mM. Even under normal conditions glutathione removal from mature erythrocytes is considerable, leading to a turnover half-time of 2–4 days [68]. This export rate confronts erythrocytes with a problem. They are not able to import glutathione directly over the plasma membrane. Moreover, mature erythrocytes lack γ -glutamyltranspeptidase activity [137, 138]. As a result of this, erythrocytes are not able to recover the glutamic acid or glycine moiety of GSH from the S-conjugates, but have to excrete the complete GSH conjugate. γ GT is also an important enzyme in the γ -glutamyl cycle [40, 96]. The absence of γ GT from erythrocytes therefore also means that they cannot utilize γ -glutamyl cycle activity for efficient amino acid uptake.

Synthesis of glutathione in erythrocytes is catalysed by γ -glutamylcysteine synthetase [eq. 2.3] and glutathione synthetase [eq. 2.4] [90]. The activity of γ -



Enzymatic synthesis of glutathione is catalysed by γ -glutamylcysteine synthetase [eq. 2.3] and glutathione synthetase [eq. 2.4].

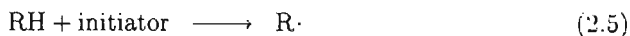
glutamylcysteine synthetase in human erythrocytes is quite low (about $0.25 \text{ U} \cdot \text{g Hb}^{-1}$ [19]), but would still be sufficient for a glutathione turnover half-time of less than 20 min. γ -Glutamylcysteine synthetase is feedback inhibited by glutathione, and this is probably a crucial factor in regulation of erythrocyte glutathione levels [119]. Under normal conditions the two enzymes are nearly saturated with ATP [19], since the K_m values of approximately 0.4 mM for both enzymes [110] are well below the ATP concentration of 1.4 mM found in normal erythrocytes. Substrate availability, mainly of glutamate, is considered to be a serious limiting factor in erythrocyte glutathione synthesis [19]. The permeability of the erythrocyte membrane for glutamic acid is very low [47], and increased plasma glutamate levels do not lead to increases in erythrocytes [139]. Several other

sources for glutamate in erythrocytes have been suggested. Direct conversion of glutamine to glutamate, as suggested by Ellory and coworkers [47], is not possible since erythrocytes do not contain glutaminase activity [79,122]. The amino group of glutamine can also be transferred to nicotinic acid [115] as part of the NAD synthesis. It is unlikely however that NAD synthesis in erythrocytes is sufficiently high to provide the amounts of glutamate needed for glutathione synthesis. King and Kuchel [79] suggested that uptake of (α)-glutamate containing di- and tripeptides and subsequent cleavage could supply the glutamate. Again this is unlikely because of the expected low concentrations of these peptide fragments in plasma [85]. Glutamate can also be produced from oxaloacetate and aspartate by glutamic oxaloacetic acid transaminase activity. The practical importance of this route remains unclear [19]. It can be concluded from the above that limitation of glutamate availability is likely to restrict the erythrocyte capacity to synthesize glutathione under conditions where more than normal amounts of glutathione are lost. It is also possible that cysteine availability becomes limiting under some conditions. Both cysteine and glycine are imported by sodium dependent uptake systems [47,120]. In sheep a hereditary disorder has been described where erythrocyte glutathione deficiency may be caused by inadequate cysteine transport [166].

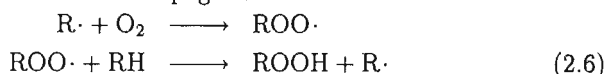
Oxidative Stress in Erythrocytes

The human erythrocyte is a cell that one might expect to be at risk for free radical damage [27,34,63]. It is exposed to high oxygen tension, especially during passage through the lungs. Erythrocytes are rich in polyunsaturated lipids [42] and iron [66]. Methyl groups between two cis double bonds in polyunsaturated lipids—as they exist in for instance arachidonic acid—are especially sensitive to lipid peroxidation [27,58,128]. The radical formed after hydrogen abstraction from such a methyl group is resonance stabilized by, energetically favorable, conjugated dienes. After reaction with oxygen and abstraction of a hydrogen atom from somewhere else—most often from another polyunsaturated lipid, resulting in a chain reaction of lipid radicals—a lipid peroxide molecule is formed (See eqs 2.5 to 2.7). Free iron and free cupric ions can catalyse reaction 2.8 (the Fenton reaction), in which the potent hydroxyl radical ($\text{OH}\cdot$) is formed from hydrogen peroxide (H_2O_2). To complete the Haber-Weiss cycle the Fe^{3+} has to be reduced by either hydrogen peroxide [eq. 2.9] or superoxide [eq. 2.10]. Apart from this, free iron can also stimulate hemoglobin autooxidation and stimulate the decomposition of hydroperoxides (see below for those two aspects). During erythrocyte life there is a constant conversion of oxyhemoglobin to methemoglobin. Normally about 3% of the total hemoglobin is converted to methemoglobin each day [25]. This methemoglobin is reduced by NADH methemoglobin reductase, which depends on the glycolytic pathway for NADH reduction [155]. An NADPH depen-

Initiation



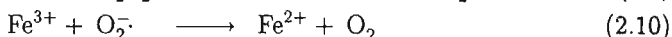
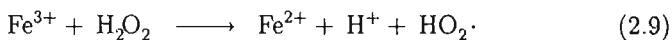
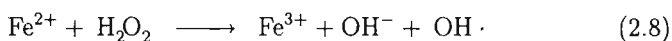
Propagation



Termination

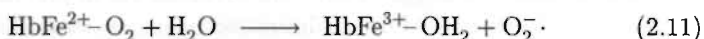


Classical description of the three phases in free radical reactions. In the initiation phase [eq. 2.5] a free radical is formed. During the repetitive propagation phase [eqs. 2.6] an existing radical will, in most cases, first react with oxygen. Next, the peroxy radical formed abstracts a hydrogen atom from another molecule (most often a polyunsaturated fatty acid), yielding a new radical. Finally, it is conceivable —although not likely— that two free radicals meet each other. This would then lead to termination [eqs. 2.7].



The Fenton reaction [eq. 2.8] leading to formation of hydroxyl radicals from hydrogen peroxide. The Fe^{3+} formed can be reduced by either another hydrogen peroxide molecule or by a superoxide radical. In this way superoxide radicals could stimulate the formation of more potent hydroxyl radicals [13, 121].

dent methemoglobin reductase —depending on the hexose monophosphate shunt for NADPH reduction— is also present in human erythrocytes. Under normal conditions the activity of this enzyme accounts for only about 20% of total methemoglobin reduction [167], but it may be of much more importance during stress conditions [127]. The autoxidation of oxyhemoglobin involves the formation of a superoxide radical [25, 102, 165]. Normally the heme iron electron binding O_2 returns to the iron when oxygen is released, and the iron remains in its ferrous Fe^{2+} state. When water or other small anions enter the heme pocket —which is normally prevented due to the hydrophobic nature of this pocket— oxygen can be displaced with an extra electron, i.e. as superoxide [eq. 2.11]. The autoxidation of oxyhemoglobin can be further increased by Cu^{2+} and Fe^{3+} ions present



Superoxide radical ($\text{O}_2^{\cdot-}$) formation during autoxidation of oxyhemoglobin. See [25] for a schematic representation of the mechanism involved.

in the red cell [25,121], and this also leads to superoxide radical generation [25]. As a result of this phenomenon there is a continual spontaneous source of activated oxygen within the erythrocyte. Methemoglobin forming substances that stimulate this mechanism, for instance because they form relatively hydrophobic anions that can more easily enter the heme pocket, will also stimulate superoxide radical formation. The results presented in chapter 5 indicate that this may be the case for hydroxylamines (see also [140]).

Apart from this internal source for active oxygen species, erythrocytes circulate in an environment in which they are exposed to extracellular sources of free radicals. Granulocytes and macrophages can generate hydrogen peroxide and superoxide anions that can damage the erythrocytes [28,29,162]. This formation of active oxygen species by granulocytes and macrophages is part of the normal inflammatory activity of these cells. Mature polymorphonuclear leukocytes are capable to ingest, kill and digest microorganisms. They can be stimulated by microorganisms and a variety of chemicals to a respiratory burst. This process is characterized by increased oxygen uptake, a shift in glucose metabolism towards the hexose monophosphate shunt (yielding NADPH), activation of an NADPH oxidase and generation of a variety of toxic oxygen species including $\text{O}_2^{\cdot-}$, H_2O_2 and $\text{OH}\cdot$ [27,109]. One could argue that for this reason oxidative stress in erythrocytes might be increased during inflammatory diseases [34]. Indeed, Machiedo and coworkers describe decreased red blood cell deformability in sepsis patients [89]. Decreases in red blood cell deformability are a known effect of erythrocyte membrane peroxidation, probably resulting from protein crosslinking and lipid structure disturbances [36]. In the study by Machiedo *et al.* the red blood cell deformability was found to correlate not only to sepsis state but also to malon dialdehyde levels in plasma, supporting the theory that the deformability decreases are induced by lipid peroxidation. Supportive evidence came from a rat study where the radical scavenger α -tocopherol was found to prevent red blood cell deformability decreases after induced sepsis [113]. Interestingly, red blood cell deformability decreases in patients did not only correlate to sepsis—which could actually be predicted two days before normal diagnosis—but also to multiple-system organ failure. This is an often fatal condition which has previously been related to oxidative stress [13,34].

The oxidative stress resulting from ischemia-reperfusion episodes [13,33,34,52,95] forms another possible external source for exposure of erythrocytes to oxidative stress. In erythrocytes from patients with various kinds of circula-

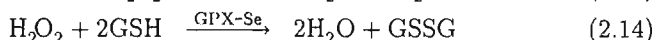
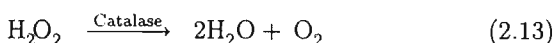
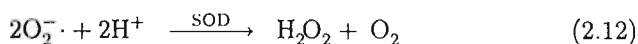
tory shock (mostly hypovolemic and cardiogenic) increased lipid peroxidation was detected by determination of aldehydes [112]. This indicates that serious hypoxia-reperfusion situations may indeed induce lipid peroxidation in erythrocytes.

From the above, it can be concluded that several factors may increase oxidative stress in human erythrocytes. Among these are increased oxygen turnover, for instance as a result of increased physical activity (studied in chapter 8 of this thesis), inflammatory diseases for instance in coal workers pneumoconiosis (studied in chapter 9) and exposure to oxidative chemicals or chemicals leading to increased methemoglobin formation (studied *in vitro* in chapter 5).

Glutathione Protection against Oxidative Stress

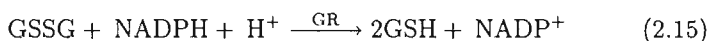
In order to understand the role of glutathione in protection against oxidative stress and more specifically lipid peroxidation we have to focus on some aspects of the oxidative processes described, and to look at the protective mechanisms that are present within the erythrocyte. The primary initiation of the radical chain-reaction is normally the formation of an active oxygen species, for instance during hemoglobin autoxidation, neutrophil activation or in the Fenton reaction (see above). When such active oxygen species reach the lipid membrane they can abstract a hydrogen atom from a polyunsaturated fatty acid. The exact nature of the actual active oxygen species involved remains unknown, but the hydroxyl radical is considered to be a likely candidate [121,128]. Due to its high reactivity ($\sim 10^9 \text{ mol}^{-1}\text{sec}^{-1}$) it will react with the nearest target available. The predominant reaction of poly unsaturated lipids observed can however also occur during the subsequent propagative chain reactions. The hydrogen abstraction by an active oxygen species is often called the initiation step in lipid peroxidation, strictly speaking it should be denominated a propagation reaction [58]. Once a lipid radical is formed a chain reaction of lipid peroxidation can commence, until either two radicals meet in a termination reaction or a stable (e.g. α -tocopheryl) radical is formed.

The erythrocyte contains protective systems that can detoxify the active oxygen species formed —thereby preventing lipid peroxidation— and that can terminate the peroxidation process. Three enzymes are present that detoxify active oxygen species: superoxide dismutase, catalase and glutathione peroxidase. The superoxide dismutase reaction [eq. 2.12] results in the formation of H_2O_2 . The longevity of H_2O_2 , its ability to pass through lipid bilayers and its reactivity towards iron that will produce the highly reactive hydroxyl radical [eqs. 2.8 to 2.10] make H_2O_2 an extremely dangerous molecule [9,126]. Its formation by superoxide dismutase is therefore in itself not necessarily a detoxification. H_2O_2 itself can be detoxified by either catalase [eq. 2.13] or the selenium dependent glutathione peroxidase [eq. 2.14]. From studies with resealed, normal and acatalasemic, erythrocytes overloaded with either catalase or glutathione [126], it can



Reaction equations for the three main antioxidant enzymes present in erythrocytes. Superoxide radicals can be converted to hydrogen peroxide by superoxide dismutase (SOD: eq. 2.12). The hydrogen peroxide itself has to be detoxified by either catalase [eq. 2.13] or selenium dependent glutathione peroxidase [eq. 2.14].

be concluded that catalase is likely to be the most important of the two. The opposite view has been presented in older literature though [31]. The glutathione peroxidase reaction with H_2O_2 [eq. 2.14] results in the oxidation of GSH. When sufficient NADPH is available this will be reduced by glutathione reductase [87], a reaction in which NADPH is oxidized to NADP^+ [eq. 2.15]. The NADPH can

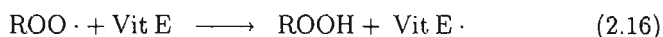


By means of the glutathione reductase (GR) reaction the erythrocyte is able to reduce glutathione. The reaction consumes NADPH that has to be provided by the hexose monophosphate pathway.

normally be regenerated by the hexose monophosphate shunt, which has a high reserve capacity. Under oxidative stress conditions up to 92% of the phosphorylated glucose can pass through this pathway —normally about 11%— [4]. Under such conditions the turnover is no longer limited by glucose 6-phosphate dehydrogenase but by phosphorylation of glucose itself, catalysed by hexokinase [153]. It was proved by Llobell *et al.* [86] that, at least in yeast, the activation of glucose 6-phosphate dehydrogenase by GSSG is actually a result of NADP^+ formation by glutathione reductase. Using this knowledge it is possible to discriminate hemolytic drugs that directly consume NADPH (like methylene blue) from those that primarily oxidize GSH (like doxorubicine) [69]. Under high oxidative stress conditions glutathione reductase activity may become insufficient and toxic amounts of GSSG have to be removed in another way. As we shall see in the next few paragraphs, oxidation of GSH by glutathione peroxidase activity towards H_2O_2 will most likely be only a small fraction of the total glutathione oxidation under oxidative stress conditions. NADPH is also of importance for catalase activity —albeit as a cofactor and not as a substrate— and it was suggested by Scott and coworkers that this might be one of the reasons why NADPH, and not GSH, modulates the oxidant sensitivity in normal and glucose 6-phosphate deficient erythrocytes [127].

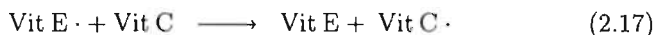
Apart from the antioxidant enzymes, the red cell contains some soluble antioxidant like ascorbate (vitamin C) and uric acid that can stabilize the active oxygen before it reaches the lipid structure [106]. Whether GSH can function as such a direct radical scavenger is debatable [106]. Such an activity would result in the formation of thiyl (GS^-) radicals, that can be stabilized by ascorbate [58, 106]. Alternatively a thiyl radical might react sequentially with another GSH and oxygen and yield GSSG and O_2^- [106].

Once lipid peroxidation has been initiated the most important protective factor is tocopherol [106, 107, 151]. When a lipid peroxy radical reacts with vi-



Termination of lipid peroxidation by formation of a stable chromanoxyl radical from vitamin E.

tamin E [eq. 2.16] a fairly stable vitamin E radical is formed, as a result of electron delocalisation in the chromanoxyl head. The concentration of vitamin E in membranes is relatively low [32], and it has to be recycled in order to obtain the protection seen in erythrocytes. Already in the original studies by Tappel [151] it was shown that vitamin E radicals can be restored by vitamin C. Recently elegant studies confirmed a direct interaction between the two [16, 32]. The ascorbyl radical thus formed can be enzymatically reduced back to vitamin



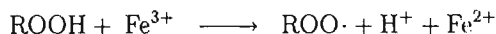
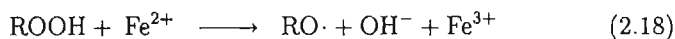
Restoration of vitamin E from vitamin E radicals by vitamin C.

C by NADH-dependent enzymes [27]. It can also react further with a second tocopheryl radical yielding dehydroascorbate [16], or two vitamin C radicals can disproportionate yielding an ascorbate and a dehydroascorbate molecule [20]. The formation of dehydroascorbate by either of those two mechanisms is of special interest to us, since dehydroascorbate can be reduced by glutathione, a reaction that may be catalysed by thioltransferases [164]. In this way glutathione could assist the vitamin E radical scavenging. The existence of a free radical reductase [94], whereby GSH could directly reduce vitamin E radicals, was suggested by the work of Reddy *et al.* [118], and further evidence for its existence was presented by Haenen *et al.* [56, 57, 58]. Fuji and coworkers also described a heat-labile glutathione dependent protection system in erythrocytes that was not identical to glutathione peroxidase or glutathione S-transferase [54]. It can be speculated that they were also looking at free radical reductase activity. Both the direct free radical reductase activity and the dehydroascorbate reduction would result

in increased glutathione oxidation. Thereby adding to the glutathione oxidation rate resulting from hydrogen peroxide peroxidase activity as described above.

So far the role of glutathione in the protection against reactive oxygen species in the aqueous phase and in the termination of lipid peroxidation were discussed. Two other important protective functions of glutathione remain to be described: the removal of lipid hydroperoxides and the protection of protein thiols.

Lipid peroxidation can not only be initiated by reactive oxygen species (the so called lipid hydroperoxide independent lipid peroxidation). It is also possible that a lipid hydroperoxide, formed as the result of another reaction chain, decomposes [eqs 2.18 and 2.19]. As a result of this new lipid radicals are formed, which will result in a whole new propagating lipid peroxidation chain reaction (i.e. lipid hydroperoxide dependent lipid peroxidation). Two separate mechanisms for this

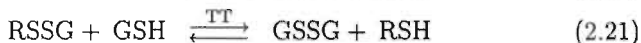


Amplification of lipid peroxidation by iron catalysed [eq. 2.18] bimolecular and hemoglobin induced unimolecular [eq. 2.19] decomposition of lipid hydroperoxides.

phenomenon are described in the literature. Metal-catalysed decomposition depends on free metal ions (e.g. iron ions in eqs. 2.18 [128]). The bond dissociation energy for a simple alkyl hydroperoxide is so high ($\sim 47 \text{ kCal mol}^{-1}$) that spontaneous unimolecular decomposition [eq. 2.19] can be excluded [27]. Hemoglobin will however catalyse this activity, a close look at the mechanism might however lead to something very close to equation 2.18. One of the main protective functions of glutathione is probably that it can prevent this amplification of lipid peroxidation by detoxification of the lipid hydroperoxides themselves [see eq. 2.1 on page 18]. Erythrocytes contain both a hydrophilic and a hydrophobic form of the selenium dependent glutathione peroxidase [51]. The hydrophobic form can directly interact with phospholipid hydroperoxides. Before the hydrophilic form can become active the lipid hydroperoxide must be removed from the glycerol moiety by phospholipase activity [82, 150]. The same is true for the glutathione peroxidase activity of glutathione S-transferase. The activity of all three enzymes—i.e. selenium dependent hydrophobic, selenium dependent hydrophilic and glutathione S-transferase—activity will lead to the glutathione oxidation.

Finally, glutathione can be used to reduce protein mixed disulfide [74], or to reduce cystine bridges. An example of the first type of reactions is given in equation 2.2 on page 18. The reduction of a cystine bridge would be a two

step process [164], in which the disulfide bridge is reduced at the expense of



Thioltransferase (TT) catalysed reduction of a disulfide bridge, e.g. an intramolecular cystine bridge.

glutathione oxidation. In human erythrocytes this type of reactions are catalysed by thioltransferase [99,100].

It can be concluded that all four protective mechanisms described —i.e. hydrogen peroxide peroxidase activity, reduction of vitamins C and E, lipid hydroperoxide peroxidase activity and thioltransferase activity— will lead to glutathione oxidation. Under serious stress conditions glutathione reductase activity may become insufficient to hold the high GSH/GSSG ratio needed for optimum protection. The GSSG export system described on page 19 will then remove part of the GSSG and even under less severe stress conditions this export system may compete with glutathione reductase. Under severe stress conditions the GSSG level may still become high enough to allow formation of mixed protein-glutathione disulfides by the reversal of the thioltransferase catalysed reaction shown in eq. 2.2. Since GSSG export is ATP dependent, insufficient supply of ATP might also stimulate this pathway. Apart from the studies present in this thesis some other reports have become available that show that human erythrocyte GSH may be a valuable marker for oxidative stress. Sugawara and coworkers found increased lipoperoxide concentrations, lowered superoxide dismutase and catalase activities and lowered glutathione concentrations in workers exposed to lead [143]. On the other hand, Saralakumari and Rao showed that in chronic fluoride poisoning —leading to oxidative stress and increased lipid peroxidation— erythrocyte GSH was increased two-fold [123]. They also reported increases in γ -glutamylcysteine synthetase, glutathione S-transferase, glutathione peroxidase and —hard to believe— γ -glutamyl transpeptidase, and concluded that an adaptation to the permanent stress must have occurred.

Glutathione S-Transferase

The main glutathione S-transferase present in human erythrocytes —often denoted GST ρ [93]— belongs to the π class [21,158]. This isoenzyme accounts for 90–95% of the total erythrocyte GST activity (using the common substrate CDNB (1-chloro-2,4-dinitrobenzene) [19,50]. The first 20 amino acids of the amino terminal sequence are the same as for GST3 (or GSTP1-1 as it should be called according to new nomenclature [92]), the π class isoenzyme from human

placenta. While the two were not discriminated immunologically, the erythrocyte isoenzyme has a higher isoelectric point [146]. Whether this difference is the result of differences in genetic origin or of post-translational modifications is difficult to evaluate. It may not even be unique for the erythrocyte since structural analysis of human placenta GST also demonstrated coexpression of at least two GST π genes [1].

Apart from GST ρ , erythrocytes contain another GST with a much higher isoelectric point ($pI > 10$ as opposed to $pI \approx 4.5$ [93] for GST ρ), which has been designated GST σ [10]. It can be separated from the major ρ form by affinity chromatography on bromosulphophthalein-S-glutathione columns. This minor isoenzyme has an amino acid content different from the ρ form and shows no immunological cross reactivity. Contrary to GST ρ it will catalyse GSH conjugation of bromosulphophthalein and 1,2-epoxy-3-(p-nitrophenoxy) propane [133]. GST σ represents approximately 5% of the total erythrocyte activity towards CDNB. Its importance might be underestimated however, since it has a comparatively low affinity to this common substrate [133].

More recently another erythrocyte GST, without activity to CDNB, has been described [60,124]. This GST is important for the conjugation of methylene chloride. It was separated from GST ρ and GST σ by affinity chromatography on GSH sepharose. Based on inhibition studies with deoxycholate and N-ethyl maleate it was further discriminated from μ class GSTs. The absence of affinity to GSH sepharose and the lack of activity towards CDNB explain why this GST was not described earlier. Schröder *et al.* [124] concluded that the newly discovered erythrocyte GST shows some resemblance to GST θ . The Theta class is the most recently described class of cytosolic GST isoenzymes —after Alpha, Mu and Pi— apart from human liver GST θ it contains the rat liver isoenzymes GST 5-5 and 12-12 [98], and the rat liver mitochondrial GST 13-13 [61]. The GST activity towards methylene chloride and some other C1/C2 compounds (methyl bromide, methyl iodide [59], dichloromethane [152] and ethylene oxide [53]) in human erythrocytes shows polymorphism [60]. This might explain the findings by van Doorn *et al.* [43], who described that four workers exposed to methyl chloride excreted high levels of a thioether metabolite, while two others exposed under the same conditions excreted almost nothing of this metabolite. This polymorphism is not only of importance for biological monitoring, but also for risk evaluations. This was confirmed in the study by Hallier *et al.*, who found that the induction of sister chromatid exchanges by methyl bromide, ethylene oxide and dichloromethane was much lower in blood of so called “conjugators” (i.e. individuals with high activity of the described GST) than in that of “non-conjugators” [60]. Risk evaluations are hampered further by the finding that the polymorphic presence of a C1/C2 conjugating GST seems to be limited to human erythrocytes [41]. In the studies described in this thesis GST activities were determined using CDNB, while protein levels were sometimes determined using antibodies to hGSTP1-1. Both assays do not pick up the polymorphic GST. A small fraction of the total

detected GST activity (about 5%) will originate from GST σ . Western blots of isoelectric focusing gels proved that it is not detected by the antibody used in chapter 6.

Normal erythrocyte GST activities towards CDNB vary considerably between individuals (6–10 fold) [125, 141] (also confirmed in this thesis), and it was suggested early that the cause might be undetected (genetic) variation [142]. Fazi and coworkers showed that GST ρ actually exists in two forms [50] that can be separated using ion-exchange chromatography. Only one isozymic form is present in young erythrocytes (separated from older cells by using density gradients), while older erythrocytes contain two forms, with up to 26% in the “secondary” form. The “secondary” GST ρ (i.e. the form that is only present in older cells) is more heat stable than the “primary” form. The kinetic properties of the two isozymic forms (K_m for CDNB and GSH, pH optimum and inhibition by 1,2-dichloro-4-nitrobenzene, 1,2-epoxy-3-(p-nitrophenoxy) propane and bromosulphophthalein) do not differ. In the same study a decrease in GST activity with erythrocyte age was also demonstrated. Combination of the two results indicates that the activity (V_{max}) is likely to be lower for the “secondary” form, but this was not actually measured.

Another interesting study towards interindividual differences in human erythrocyte GST activities was reported by Anosike and coworkers from Nigeria [7]. They compared GST activities in erythrocytes from 20 HbSS individuals—homozygotes for HbS, the sickle cell form of Hb—, with those for 13 HbAS—HbA, HbS heterozygotes— and 20 HbAA—normal— individuals. GST activities were found to be increased in HbS individuals. Mean values were 14.61 U·g Hb⁻¹ for HbSS, 4.65 U·g Hb⁻¹ for HbAS and 2.75 U·g Hb⁻¹ for HbAA. The sickle cell disease is characterized by hemolytic crisis which cause anemia and other subsequent complications. The nature of those hemolytic crisis is not fully understood, but they are associated with methemoglobin formation, superoxide release and liberation of heme. Van den Berg *et al.* [15] have shown that the lipid peroxidation promoting capabilities of HbS (and heme) are higher than that of normal Hb. When heme is liberated it will further increase the damage to the erythrocyte, as heme itself has been shown to cause hemolysis *in vitro* [80]. Erythrocyte GST has a high affinity towards heme, and it was suggested by Harvey and Beutler that one of the purposes of the enzyme is to transport newly synthesized heme from mitochondria to be incorporated into globin [62]. It is tempting to speculate—as was done by Anosike *et al.*— that the extra GST present in HbS subjects is synthesized during hematopoiesis to protect the cell against the oxidant species formed or to scavenge heme or both.

Beutler *et al.* [18] found GST deficiencies in two sibs with glutathione synthetase deficiency and GSH levels of less than 5% of normal. Since the parents had half-normal glutathione synthetase activities and normal GST, the authors concluded that deficiency in glutathione synthetase was the primary lesion and that GST appeared to be labile at the very low glutathione concentrations that re-

sulted thereof. Two different explanations can be given for this phenomenon. The explanation given by Beutler *et al.* is that binding of GSH may directly stabilize GST. The K_m for GSH binding is about 0.1 mM for most soluble GSTs [91] and for the human erythrocyte form values of 0.14 mM [7] up to 0.58 mM [50] have been reported. Since physiological erythrocyte GSH concentrations are about 2 mM this means that GST will normally be saturated with GSH. This leads to conformational changes that may stabilize the enzyme, as it is known to do *in vitro*. Decrease of the erythrocyte GSH concentration to 5% or less of the normal 2 mM means that its value comes below the GST K_m for GSH, and that a large fraction of the GST will be in its more labile non-GSH form. The other possibility is that GSH protects GST in its soluble form. The protective mechanisms of GSH described above might also function to protect GST ρ . More specifically, GSH might assist in the prevention of lipid peroxidation —thereby also protecting GST for reactive peroxidation products— and, as a substrate of thioltransferase, GSH might help to restore GST activity when lost as a result of disulfide formation.

That the latter mechanism might actually work was shown as a side effect of the work done by Shen *et al.* [130]. Rat GST-P and human GST π (currently named rGSTP1-1 and hGSTP1-1 [92]) are known to contain 4 cysteine residues at corresponding positions (i.e. at the 14th, 47th, 101th and 169th position) [91]. The activity of these π class enzymes were found to be vulnerable to N-ethylmaleimide, a well-known SH modifier [149]. Tryptic digestion was used to identify the cysteine at position 47 as the critical target. When hydrogen peroxide treatment was used these GSTs were also inactivated, but they could be reactivated with purified rat liver thioltransferase [130]. The positions of the cysteinyl residues were identified using recombinant mutant GST-P forms with alanine instead of one of the four cysteine residues [131]. Again the cysteine 47 residue was found to be critical, but cysteine 101 was also important. Disulfide formation between these residues, resulting in steric hindrance, was considered the most probable cause for the inactivation. It may be noteworthy that, while the elegant work of this Japanese group shows that cysteine residues 47 and 101 are probably located at or near the active center of GST-P, it also shows that the cysteinyl nature of these residues is not critical. After all the two mutant forms with alanine instead of the cysteines at these locations as well as the mouse GST II, which contains glycine instead of the cysteine at location 101, all do show GST activity [131].

The possibility should be considered that the “secondary” GST ρ form described by Fazi *et al.* [50] is actually a form in which one or more of the sulphydryl groups is masked, like the S-modified forms reported by Tamai *et al.* [149], or the S-oxidized forms reported by Shen *et al.* [130].

The inactivation by oxidative stress seems to be more or less specific for class π GSTs. Nishinaka *et al.* [108] showed that class π porcine lens GST could be inactivated by biological disulfides, by a xanthine-xanthine oxidase system, by H₂O₂ and by the cataract inducing oxidant 1,2-naphthoquinone. On the other

hand, bovine lens GST—which belongs to class μ —was not sensitive to these factors. In fact, class μ GST as well as microsomal GST, can even be activated by active oxygen species [6, 88, 104]. GSTs from the α class can be inactivated by AOS [104], but they are less sensitive than π class GST.

Apart from model sulfhydryl reagents, like the N-ethyl maleimide used by Tamai *et al.* [149], several other substances are known to inhibit erythrocyte GST *in vitro*. Singh and Awasthi showed for instance that 1.3 mM of the herbicide 2,4-D (2,4-dichlorophenoxyacetate) suffices to inhibit 50% of purified GST ρ and GST σ activity towards CDNB [132]. For 2,4,5-T (2,4,5-trichlorophenoxyacetate) these I50 values are 0.5 mM and 1.2 mM for GSTs ρ and σ respectively. Ansari and coworkers showed that the industrially important chemicals acrolein, propylene oxide, styrene oxide, ethylene dibromide and ethylene dichloride also inhibited GST from erythrocytes both *in situ* and in purified GST ρ [8]. They were the first to suggest the use of human erythrocyte GST activity as a biological marker for chemical exposure. Before that, Kilpikari and Savolainen [78] had measured GST activity in erythrocytes of workers exposed to hot rubber fumes. The rationale for these measurements was the expectation that the high production of thioethers found in such workers might lead to an induction of GST activity. Their finding of a *decrease* in GST activity was therefore opposite to their expectation and not further investigated. It can be concluded that various indications do exist—some of which were only published while the work in this thesis was in progress—to make a further study towards changes in human erythrocyte GST during electrophilic and oxidative stress conditions worthwhile.

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Chapter 3

Influence of Glutathione on the Formation of Cysteine Alkylation Products in Human Hemoglobin

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Published in: *Toxicology* 52, 1988, 177–186.

Summary. Human blood samples were treated *in vitro* with iodoacetamide. At low concentrations — less than 1 mM — only a low fraction of the β -93 cysteine in hemoglobin was alkylated, whereas the alkylating reaction with glutathione was extensive. At higher iodoacetamide concentrations the glutathione pool became exhausted leading to more than proportional increases in the alkylation of the sulfhydryl group in hemoglobin. When diethyl maleate was used as a glutathione depletor prior to incubation with iodoacetamide, low concentrations of iodoacetamide were sufficient to obtain high degrees of hemoglobin sulfhydryl alkylation. N-Ethylmaleimide could not be used as glutathione depletor because the reaction with glutathione appeared to be reversible. The lower reactivity of the thiol group in hemoglobin in comparison with that of glutathione was also found for the isolated biomolecules. The protection of hemoglobin by glutathione present in the human erythrocyte renders the measurement of hemoglobin alkylation less attractive for biological effect monitoring. The sensitivity of such methods is lowered, while the important relation between the alkylation of hemoglobin and that of DNA in the target tissues is affected by interindividual differences in the ratios of the effectiveness of glutathione protection between erythrocytes and target cells.

Introduction

Many genotoxic chemicals are alkylating agents able to react with biological macromolecules. Their reaction with certain nucleophilic centers in DNA is the most probable cause of their genotoxic activity (Miller and Miller, 1966). Human exposure to these chemicals or to their inactive precursors may give rise to unwanted effects, like an increased risk of cancer. Determination of protein or nucleic acid alkylation products is currently considered as one of the most promising methods for monitoring human exposure to genotoxic chemicals (Leonard *et al.* 1987; Farmer *et al.* 1987). Because DNA (rather than protein) is believed to be the primary target of genotoxicity, the determination of protein alkylation products provides only an indirect estimation of the biological effective dose. Nevertheless, the blood proteins hemoglobin and albumin are often chosen for molecular dosimetry because they can be easily obtained while especially hemoglobin has a long lifetime (about 120 days; compared to 20–24 for albumin). A relatively low turnover rate has the advantage that it may provide an integrated index of exposure. Studies on hemoglobin alkylation may be complicated by the fact that hemoglobin is protected against chemically reactive agents by the erythrocyte membrane and by the presence of detoxifying enzyme systems in the erythrocyte. In a previous study (Evelo *et al.* 1987) we showed that hemoglobin is more effectively alkylated at the same dose of alkylating reagent after purification than in whole blood. In the work reported here we studied the role of glutathione in the protection of hemoglobin.

Materials and Methods

Chemicals

Diethyl maleate (DEM), 5,5'-dithio-bis(2-nitrobenzoic acid) (DTNB), 4,4'-dithiodipyridine (4-PDS), iodoacetamide (IAcA) and N-ethylmaleimide (NEM), were from Janssen Chimica (Beerse, Belgium), glutathione (GSH) was from Boehringer (Mannheim, F.R.G.), SephadexTM G-25 (medium) was from Pharmacia (Uppsala, Sweden). All other chemicals were of analytical purity. Only freshly deionized distilled water was used.

In Vitro Incubations

Before each experiment fresh blood samples from a single healthy volunteer were collected in vacutainers, containing ethylenediamine-tetraacetic acid (EDTA) to prevent coagulation. In a shaking waterbath (210 rev./min) closed sterile vials (I.D. 24 mm, height 55 mm) each containing 1.5 ml whole blood or 1.5 ml of a 1 mM solution of purified hemoglobin in PBS (phosphate-buffered saline; 10 mM sodium phosphate + 150 mM sodium chloride + 1 mM EDTA) or 1.5 ml 1 mM GSH in PBS were incubated at 37°C. Sterile solutions of IAcA, DEM or NEM were added in 50 μ l volumes. NEM and IAcA were dissolved in PBS, DEM in ethanol.

Isolation of Hemoglobin (Hb)

All purification steps were carried out at 4°C. To 1.5 ml blood 7.5 ml PBS of pH 8.0 was added. The samples were centrifuged at 1000 *g* for 5 min. Serum and buffy coat were removed by careful suction, and the erythrocytes were re-suspended with 8 ml PBS. After mixing, the samples were centrifuged again at 1000 *g* for 5 min. The supernatant was removed by careful suction, and a few erythrocytes were sacrificed to remove any remaining buffy coat. This washing procedure was repeated twice. The erythrocytes were then layered upon a solution of 30% sucrose, containing 0.5% Triton X-100 and 10 mM Tris (pH 9.0), and lysed by centrifugation at 900 *g* for 10 min as described by Scott (1970). To the lower part of the sucrose layer, containing the hemoglobin, 1/10 volume of 1 M sodium chloride was added, to optimize the ionic strength towards the removal of cellular debris (Riggs, 1981), and it was centrifuged once more at 10.000 *g* for 15 min. Finally, compounds of low molecular weight were removed from the hemoglobin solution with Sephadex G-25. The hemoglobin obtained in this way was approx. 85% pure when expressed on a hemoglobin/protein basis. Protein concentrations were determined by the Lowry method (Lowry *et al.* 1951).

Determination of Free Sulfhydryl Groups in Hemoglobin

Free sulfhydryl groups in hemoglobin (HbSH) were determined by the procedure of Grasseti and Murray (1967) as modified by Neis *et al.* (1984). The assay is

based on spectrophotometrical determination of the formation of 4-thiopyridine in the reaction of 4-PDS with HbSH. Purified Hb samples were diluted 1:150 with a 100 mM sodium phosphate buffer of pH 7.15 containing 1 mM EDTA. To 3 ml of the diluted Hb sample 200 μ l of a 1 mM 4-PDS solution was added. The absorbance at 324 nm was determined after 40 min incubation in the dark at room temperature. Corrections were made for the background absorptions of Hb and 4-PDS.

The concentration of Hb was calculated from the absorption of the Hb blank in the Hb sulfhydryl assay. Alternatively the hemoglobin cyanide procedure of van Kampen and Zijlstra (1961) was used. As reported previously, both methods gave identical results provided Hb was initially in its reduced state (Evelo *et al.* 1987).

Determination of Reduced Glutathione

Immediately after incubation 100 μ l blood samples were prepared for the determination of GSH, essentially as described by Anderson (1985). The samples were acidified with 500 μ l of 10 mM hydrochloric acid to reduce GSH oxidation and to inactivate γ -glutamyl transpeptidase. The erythrocytes were lysed by freezing in a dry-ice acetone bath and subsequent thawing. This procedure was repeated three times. After centrifugation at 10,000 *g* for 20 min, 500 μ l of the supernatant were deproteinised with 10% (w/v) 5-sulfosalicylic acid and centrifugated again at 10,000 *g* for 20 min.

For the GSH assay 200 μ l of the deproteinised sample and 100 μ l 6 mM DTNB in 1% sodium citrate (w/v) were added to 700 μ l of a 143 mM sodium phosphate buffer of pH 7.5 containing 6.3 mM EDTA. The amount of 5-thio-2-nitrobenzoic acid formed was determined spectrophotometrically at 412 nm. Corrections were made for background absorptions of DTNB and of the blood sample.

Results

In the figures the amounts of thiol groups present are expressed relative to the initial values. Normal HbSH values were slightly over 2 (mean: 2.20 ± 0.06) free sulfhydryl groups per tetrameric Hb molecule. The Hb concentration in the blood samples used was about 2.5 mM. The initial GSH concentrations in blood varied slightly between experiments (mean: 0.7 ± 0.1). For each incubation the HbSH and GSH values are the mean of duplicate determinations.

When solutions of 1 mM purified Hb in PBS were incubated with IAcA for 1 h alkylation was found to be substantially pH dependent (Figure 3.1 on the next page). At higher pH values the extend of alkylation was higher. Analogous results were obtained with solutions of 1 mM GSH in PBS. The reaction with GSH was more effective than with Hb. When a solution of 1 mM purified Hb in PBS of pH 7.3 was supplied with GSH before 1 h incubation with IAcA the

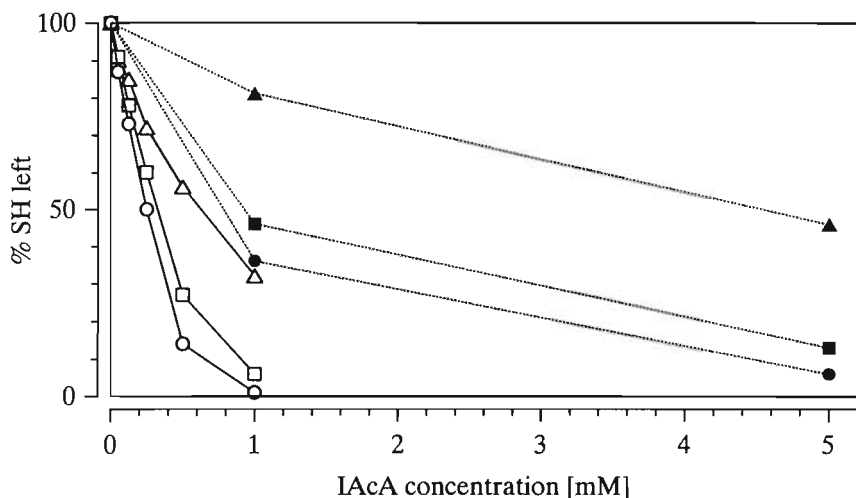


Figure 3.1: Effect of pH on the disappearance of free sulfhydryl groups from 1 mM purified Hb (.....; closed symbols) and 1.0 mM GSH (—; open symbols) after incubation with IAcA at 37°C in PBS for 1 h. Δ = pH 6.5; \square = pH 7.3; \circ = pH 8.0. Data are the mean of duplicate determinations. 100% HbSH values were 2.12, 2.27 and 2.23 respectively.

extent of SH alkylation was lowered proportional to the amount of GSH added (Table 3.1 on page 49). Whereas 33% of the HbSH was alkylated without GSH addition this percentage was lowered to 12% when 1 mM GSH was supplied, i.e. under conditions where there is one GSH thiol group present for each IAcA.

Figure 3.2 on the next page shows that in whole blood the reaction of IAcA with GSH prevailed over the reaction with HbSH. At IAcA concentrations below 1 mM hardly any reaction occurred with the Hb thiol, whereas the reaction with GSH was extensive. At higher IAcA concentrations the GSH pool was exhausted and only then the reaction with HbSH became more important.

In order to verify whether GSH really has the protective action indicated by the above results we looked for an effective GSH depletor which did not react with the SH group of Hb. Figure 3.3 on the following page shows that 15 min incubation with 2.5 mM NEM sufficed to mask nearly all GSH in whole blood. At higher NEM concentrations extensive reaction with HbSH occurs.

For DEM about 10 mM was needed to obtain the same result. For both substances the specificity was satisfactorily high. Almost 90% of the HbSH was still free under conditions where more than 90% of the GSH had been masked. In subsequent experiments the interaction between GSH and the depletors was studied in more detail. The influence of the incubation time on the disappearance of free thiol groups is shown in figure 3.4 on page 50.

In the presence of 1 mM NEM over 60% of the initial GSH concentration

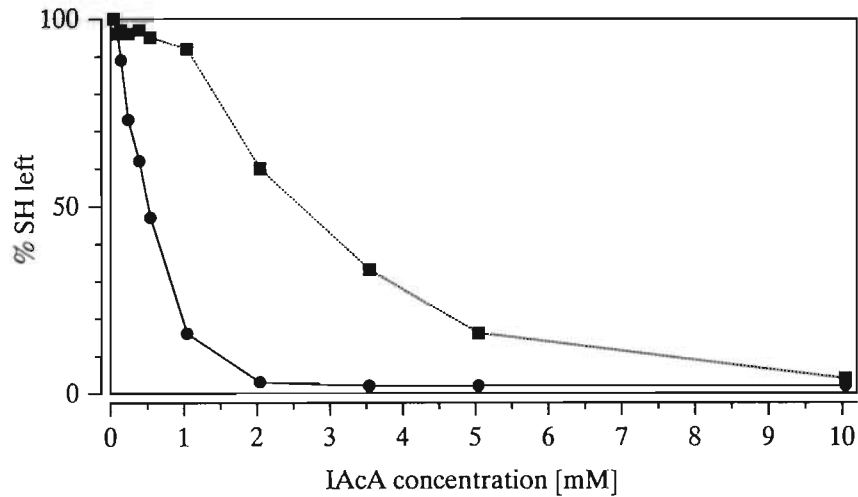


Figure 3.2: Disappearance of free sulfhydryl groups from Hb (···■···) and GSH (—●—) in blood incubated at 37°C with IAcA for 1 h. Data are the mean of duplicate determinations. The 100% HbSH value was 2.15; 100% GSH corresponds to 0.86 mM.

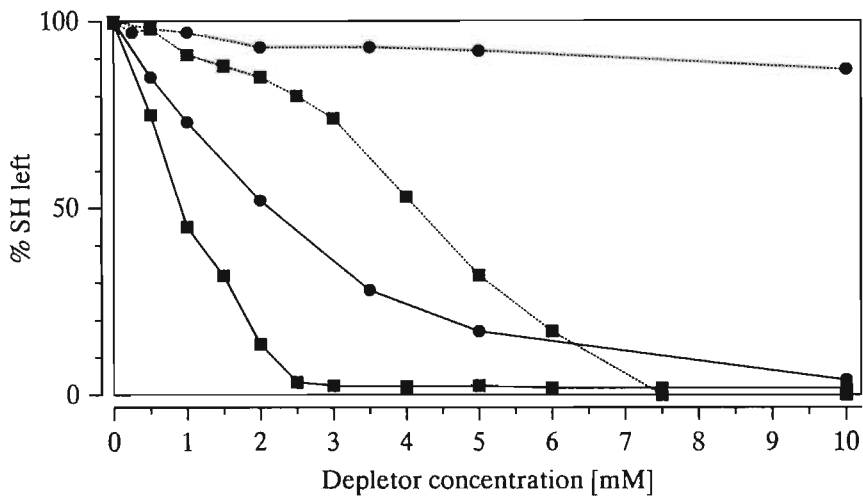


Figure 3.3: Free sulfhydryl groups of Hb (·····) and GSH (—) left after incubation of blood with the GSH depletors NEM (■) and DEM (●) at 37°C for 15 min. Data are the mean of duplicate determinations. The 100% HbSH values were 2.18 and 2.29 respectively; 100% GSH corresponds to 0.72 and 0.82 mM respectively.

GSH added mM	HbSH left [SH/Hb]
–	2.22 ¹
0.00	1.41
0.33	1.58
0.67	1.74
1.00	1.84
1.33	2.00

¹: Initial value without IAcA.

Table 3.1: Disappearance of free sulfhydryl groups from 1 mM purified Hb incubated with 1 mM IAcA at 37°C in PBS of pH 7.3 for 1 h in the presence of GSH.

was masked after 15 min incubation. Thereupon, increasing amounts of free thiols were found after longer incubation times. After 5 h incubation the concentration of free thiols had reached again 85% of the initial value. Similar incubations with 2 mM DEM were also followed for 5 h. The reaction between DEM and GSH was found to be rather slow. Under the conditions used some GSH could still be detected after more than 1 h incubation time. It was interesting to find that GSH once reacted with DEM did not become available again within the 5 h period the reaction was followed.

The reaction of IAcA with GSH and HbSH was also followed during incubation of 0.5 mM IAcA with whole blood over a five hour period (data not shown). The reaction of IAcA with both thiols was found to be fast; maximum thiol depletion was reached within 15 min. No change occurred in the amount of thiol groups detectable after this first 15 min.

Finally, blood samples were incubated with 1 mM IAcA for 1 h after preincubation with 2 mM DEM for 2 h (Figure 3.5 on the following page). In accordance with the results shown in figure 3.4 on the next page, the DEM treatment alone sufficed to mask all GSH while the majority of HbSH groups remained unreacted. As expected, the reaction of HbSH with IAcA was more pronounced after DEM treatment than without it.

Discussion

The action of the well-known GSH depletors diethyl maleate (DEM) and N-ethylmaleimide (NEM) was found to be different. DEM has been used *in vivo* (e.g. Yang *et al.* 1987) and with isolated hepatocytes (Högberg and Kristoferson

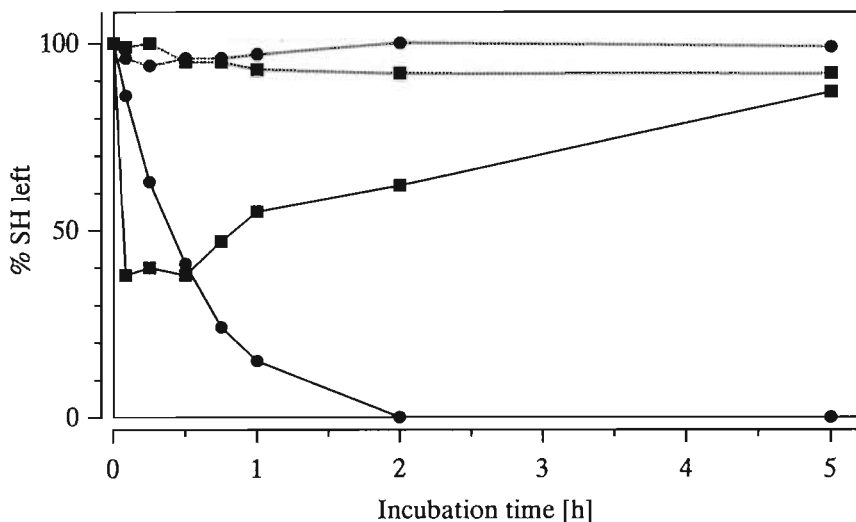


Figure 3.4: Effect of incubation time on the disappearance of free sulphydryl groups from Hb (.....) and GSH (—) after incubation of blood with 1 mM NEM (■) and 2 mM DEM (●). Data are the mean of duplicate determinations. The initial HbSH values are 2.23 and 2.30; while the initial GSH values are 0.57 and 0.61 mM respectively.

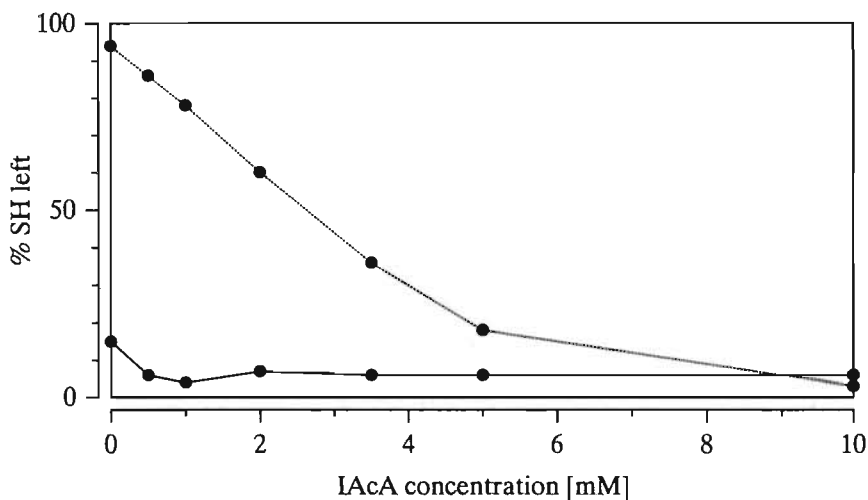


Figure 3.5: Disappearance of free sulphydryl groups from Hb (.....) and GSH (—) in blood incubated at 37°C with IAcA for 1 h after preincubation with 2 mM DEM for 2 h. Data are the mean of duplicate determinations. The 100% HbSH value was 2.19 and the original GSH value (i.e. before preincubation with DEM) was 0.72 mM.

1977). NEM is used to mask reduced GSH before the determination of oxidized glutathione (GSSG) (Tietze 1969). At the low doses needed to obtain specific masking of GSH without affecting the hemoglobin thiol the action of DEM was found to be slow but irreversible. We also found the same slow reaction in *in vivo* experiments with rats (unpublished). The reaction of NEM was much faster, but the reappearance of free thiol groups during prolonged incubation times indicated that the reaction between NEM and GSH is reversible. We concluded that for the purpose of depleting GSH from whole blood DEM is more useful than NEM. The reversibility of the reaction of NEM with GSH may have some consequences for the application of NEM in the assay of oxidized glutathione as was described by Tietze (1969). Release of GSH during that assay would lead to an overestimation of GSSG, while liberated NEM would disturb the assay because of its ability to inhibit glutathione reductase activity.

The reactivity of the model alkylating agent iodoacetamide towards the thiol group of glutathione was found to be much higher than its reactivity towards the hemoglobin thiol group. An effective protection of HbSH by GSH was also found for NEM. The difference in reactivity was not only found in whole blood but also for (the combination of) the isolated components. Therefore, it has to be attributed to differences in direct alkylating activity rather than to glutathione transferase activity. This is in agreement with the very high reactivity towards thiol groups of iodoacetamide. In the case of other, less reactive compounds activity of glutathione transferase is expected to be rate-limiting.

The conformation of the hemoglobin molecule probably lowers the reactivity of the β -93 thiol group, as it does in extreme for the other two thiol groups, which are only reactive after denaturation of the hemoglobin.

It has been suggested that the alkylation of target nucleophilic centers in DNA may be calculated from the degree of alkylation of other nucleophilic centers in proteins, using the Swain-Scott relationship (e.g. Ehrenberg 1980). The present results show that the nucleophilic strength of centers in biomolecules cannot be derived from the known values for the isolated centers but have to be carefully determined. When conformational effects are the main cause of the lowered activity the apparent nucleophilic strength of the protein centers may even be influenced by the chemical nature (e.g. their lipophilicity) of the alkylating agents.

In animal studies with various types of radioactively labeled genotoxic compounds the ratio of DNA alkylation product formation in blood cells and binding to hemoglobin was found to be constant over a large dose range (Neumann 1980). The constancy of the ratio of hemoglobin alkylation product formation and the formation of DNA alkylation products in the target tissues, not only at different doses but also between individuals, is an important prerequisite for the estimation of DNA binding on the basis of protein binding measurements. Our results suggest that interindividual differences in the ratio between erythrocyte and target cell glutathione levels (and possibly in glutathione transferase activity) will influence the ratios between erythrocyte and target dose. For individuals with a

relatively high detoxification activity in their erythrocytes, the erythrocyte dose will be relatively lower at equal target doses. As a consequence equal degrees of hemoglobin alkylation found for individuals with differing ratios of intracellular glutathione levels between erythrocytes and target cells will not correspond with equal degrees of DNA alkylation in those target cells.

The high protection of hemoglobin against alkylation found, renders this protein less promising for the purpose of biological effect monitoring. Since the plasma concentration of GSH is much lower than the erythrocyte concentration measurement of albumin alkylation is expected to be a more sensitive method.

Acknowledgment. The authors are grateful to Dr. Olthuis of the "Bloedbank Nijmegen" for the supply of the blood samples. We thank Lianne de Wijs for her technical assistance. Financial support was given by the General Directorate of Labour, Dutch Ministry of Social Affairs.

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Chapter 4

Modulation of Glutathione and Glutathione S-Transferase Levels by Cyclophosphamide and Cisplatin

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Submitted for publication.

Abstract *In vitro* modulation of glutathione (GSH) and glutathione S-transferase (GST) levels by cyclophosphamide (CP) and cisplatin was investigated in rat liver fractions and in human erythrocytes. Some antimetabolites were used for comparison. Incubations with and without a microsomal activating system (MAS) were performed. In microsomal GST studies the microsomes present were completed with a NADPH regenerating system (RS) to attain a complete MAS. Addition of RS alone (i.e. without drug addition) led to an increase in microsomal GST activity. In contrast, addition of MAS alone led to a decrease in liver cytosolic and in erythrocyte GST activity and in GSH concentrations. These effects are probably due to formation of active oxygen species by microsomes in the presence of RS. Supportive evidence for increased active oxygen species availability in erythrocytes was given by methemoglobin formation and lipid peroxidation in erythrocyte incubations with MAS.

The antimetabolites 5-fluorouracil and cytosine arabinoside did not affect the GST activity and GSH concentrations in the different incubations. Methotrexate, another antimetabolite, only inhibited the microsomal GST activity. Cisplatin, an alkylating agent, caused a GST activity decrease in incubations of microsomal and cytosolic liver fractions despite the presence of MAS. In liver cytosolic incubations GSH was depleted by cisplatin. In erythrocyte incubations with cisplatin no significant effects were seen on GST and GSH levels. This suggests that cisplatin is not able to enter the erythrocyte. In the presence of MAS, CP —another alkylating agent— gave an almost complete depletion of GSH in rat liver cytosol and in human erythrocytes. Rat liver microsomal GST, diluted rat liver cytosolic GST and GST in human erythrocytes were activated by CP in the presence of MAS. Without dilution cytosolic GST was protected by the high GSH concentration present. Without a complete MAS no significant effects for CP were found.

The modulation of GST activity and GSH concentrations by CP and cisplatin occurred during *in vitro* incubations where protein synthesis was not possible. This modulation, particularly seen here for GST class π , could be of importance in relative sensitivity of tumor tissue. Since GSH/GST will detoxify many alkylating antineoplastic agents and organic hydroperoxides formed as a result of chemotherapeutic or radiotherapeutic treatment, changes in GSH and GST levels may influence cytotoxicity and therapeutic efficacy. Therefore, selective GSH depletion in tumor cells by CP and cisplatin in combination with radiation therapy appears to be a promising approach for tumor sensitization.

Glutathione S-transferase (GST; EC 2.5.1.18) isoenzymes are found in nearly all tissues and are involved in the detoxification of a variety of noxious xenobiotics, including certain antineoplastic agents and their metabolites [1]. The best known reaction catalyzed by GSTs is the conjugation of electrophiles with the endogenous tripeptide glutathione (GSH), which results in the formation of thio-

ethers. In general, GSTs are high capacity intracellular binding proteins that, independently of their enzymatic activities, may serve in the storage, transport, or removal of many hydrophobic compounds [2,3]. Additionally, some GST isoenzymes may participate in the repair of oxidative damage to membrane lipids and DNA [3,4]. These properties indicate that GSTs function as part of an important cellular defense system against the cytotoxic effects of carcinogenic and antineoplastic agents [5].

Strong evidence exists that elevation of cellular GST and GSH levels, may occur in tumor cells in response to antineoplastic agents. This overexpression of individual GST enzymes and increases in GSH levels are associated with increased cellular resistance to antineoplastic agents [5,6,7,8,9]. Recently, much research has focussed on π class GSTs. Rat GST P (7-7) was identified as a marker for rat hepatic preneoplastic and neoplastic lesions [10]. In humans GST π may be a useful marker not only for various cancers, but also for some high-risk precancerous lesions [8]. Furthermore, absence of GST μ , another soluble GST isoenzyme, has been suggested as a possible marker for greater susceptibility to lung cancer among smokers. Hereditary differences in the expression of this form are due to deletion of the gene [11]. Apparently GSTs and GSH levels do not completely account for the resistance patterns observed in human tumors and cell lines selected for resistance to cytotoxic agents [6]. Nevertheless, further study of changes in GST and GSH levels after exposure to antineoplastic agents may lead to a better insight in the process of drug resistance in tumor cells.

GSH can protect cells against electrophiles and free radicals [12,13]. Since formation of radicals is characteristic of ionizing radiation, GSH may play an important role in radiosensitivity. Potential mechanisms of radioprotection by GSH include radical scavenging, and hydrogen donation to molecules (DNA), and reduction of peroxides and maintenance of protein thiols [13,14]. Radioprotection by GSH is not effective under aerobic conditions because the intracellular GSH concentration is not able to compete successfully with oxygen for radiation-induced radicals. Under hypoxic conditions GSH becomes more competitive and GSH depletion will have an effect on radiosensitivity. Since hypoxic cells are a characteristic feature of tumors, depletion of GSH in combination with radiation therapy can be a promising strategy for selective tumor sensitization. However, it may not always be possible to achieve selective depletion of GSH in tumors because of toxicity occurring in normal tissues [14].

In principle, changes in GSH concentrations and in GST abundance, isoenzyme expression and enzyme activity can all be used for human exposure monitoring, as long as the chosen parameter is accessible to experimental determination [15]. For practical monitoring purposes, blood is the most easily available bio-matrix containing GST and GSH. Ansari *et al.* [16] have shown that GST activity in human erythrocytes is inhibited by the industrial chemicals acrolein, propylene oxide, styrene oxide, ethylene dibromide and ethylene dichloride. Kilpikari and Savolainen [17] found a decrease in GST activity in the erythrocytes of workers

exposed to hot rubber fumes. We previously described [18] a decrease in GST activity in erythrocytes, as well as a depletion of GSH in whole blood after occupational exposure to 1,3-dichloropropene. Furthermore, GST activity in human erythrocytes and GSH concentrations in whole blood were decreased shortly after long distance running and are increased after several months of endurance training [19] and GST activities are decreased in erythrocytes of miners with an early stage of coal miners pneumoconiosis [20].

Modulation of the GSH/GST system by different antineoplastic agents was tested *in vitro* in rat liver fractions and in human erythrocytes. Erythrocytes were chosen because they contain the same class of GST isoenzyme (π class) as many tumors have in overexpression and because they are easily available. Liver fractions were chosen because this organ is susceptible to reactive metabolites due to a very high phase I metabolic activity [21] and because it has a completely different GST isoenzyme pattern (mainly α class) from that of erythrocytes (only π class) [2]. In this study five antineoplastic agents were tested which are widely used in the treatment of various cancers. Cyclophosphamide (CP) and cisplatin (c-DDP, cis-diamminedichloroplatinum(II)) are alkylating agents and therefore effects on the GSH/GST system by these two agents were expected in advance. CP is a biologically inactive nitrogen mustard analogue that requires hydroxylation by hepatic microsomal enzymes to attain its alkylating activity [22]. The rate of CP metabolism is increased in rats pretreated with phenobarbital [23]. Three antineoplastic agents belonging to the group of antimetabolites were used as controls. Cytosine arabinoside (Ara-C, cytosine β -D-arabinofuranoside), and 5-fluorouracil (5-FU) are pyrimidine antagonists and methotrexate (MTX) is a folic acid antagonist.

Materials and Methods

Analytical methods

Chemicals. 1-Chloro-2,4-dinitrobenzene (CDNB), cyclophosphamide (CP, CAS: 6055-19-2), cytosine β -D-arabinofuranoside (Ara-C, CAS: 147-94-4), 5,5'-dithio-bis-(2-nitrobenzoic acid) (DTNB), dithiothreitol (DTT), 5-fluorouracil (5-FU, CAS: 51-21-8), glucose 6-phosphate (G6P), glucose 6-phosphate dehydrogenase (G6PDH, EC 1.1.1.49), reduced glutathione (GSH), glutathione reductase (EC 1.6.4.2.), 3-hydroxyacetanilide (3-HAA, CAS: 621-42-1), NADP and the reduced form NADPH were obtained from Sigma Chemical Co. (St. Louis, MO, USA). Phenobarbital was from OPG Farma (Utrecht, NL) and 2-thiobarbituric-acid (TBA) was from Merck (Darmstadt, FRG). cis-Diamminedichloroplatinum (II) (c-DDP, CAS: 15663-27-1) and methotrexate (MTX, CAS: 59-05-2) were gifts from Multipharma B.V. (Weesp, NL). All other chemicals and solvents were of analytical quality. Only microfiltrated deionized water was used.

Animals. Male Brown Norway rats (BN/M) weighing 250–280 g were used. Housing was as previously described [24]. A non-albino strain was chosen as there are some indications that biotransformation of xenobiotics is lower in albino strains [25]. Prior to the preparation of microsomes and cytosol, phenobarbital ($1 \text{ g} \cdot \text{l}^{-1}$) was added to their drinking water for one week.

Preparation of rat liver microsomes and cytosol. The basic procedure for preparation of rat liver microsomes and cytosol was described previously [26]. To make sure that the microsomal fraction was free of any cytosolic GST and GSH, the pellet was washed with four volumes of ice-cold sodium/potassium phosphate buffer (50 mM, pH 7.4). Microsomal activating system (MAS) was composed of microsomes (final protein concentration $1 \text{ mg} \cdot \text{ml}^{-1}$) and a NADPH regenerating system (RS; 0.5 mM NADP, 5.0 mM MgCl_2 , 5.0 mM G6P and 1.0 U/ml G6PDH). In microsomal GST experiments the microsomes present were completed with RS to attain a complete MAS. The protein content in both fractions was determined according to Lowry *et al.* [27].

Preparation of erythrocytes. Human blood was obtained from the Red Cross Blood Bank "Zuid Limburg" (NL), collected in 10 ml sterile vacuum tubes containing K_3EDTA as an anticoagulant and stored at 4°C until the next day. Samples were controlled for viral infections before use. Samples of 3 to 5 persons were pooled. Erythrocytes were separated from plasma and washed three times with phosphate buffered saline (PBS; 15 mM sodium/potassium phosphate + 130 mM sodium chloride, pH 7.4) and diluted in PBS to a final hemoglobin concentration of about $100 \text{ g} \cdot \text{l}^{-1}$. The hemoglobin concentrations were determined with the hemoglobin cyanide procedure described by Van Kampen and Zijlstra [28].

Procedure. Three different GST sources were used: human erythrocytes (final hemoglobin concentration $23 \text{ g} \cdot \text{l}^{-1}$), rat liver cytosol (final protein concentration $1.7 \text{ mg} \cdot \text{ml}^{-1}$ in normal and $0.2 \text{ mg} \cdot \text{ml}^{-1}$ in diluted samples) and rat liver microsomes (final protein concentration $1.0 \text{ mg} \cdot \text{ml}^{-1}$). These sources were suspended in 100 mM sodium/potassium phosphate buffer (pH 7.4) except the erythrocyte incubations with c-DDP, which were suspended in PBS to prevent dissociation of the two chloride ligands in the extracellular matrix [29]. Incubations were performed both with and without MAS. The five drugs were used with a final concentration between 0.25 and 2 mM. The complete test system without the addition of a drug was preincubated for 5 min at 37°C in a shaking water bath (80 rpm). After that, 1 ml drug in the appropriate buffer was added and incubated for 1 h (final volume test system 3 ml). After incubation with erythrocytes, red cells were washed three times with PBS and packed by centrifugation (5 min, $2000 \times g$). When applicable GSH concentrations and GST activities were determined in the same incubations. All experiments were performed three times on different days and each day all incubations were performed in triplicate.

Determination of GST. For the determination of GST activity in erythrocytes af-

ter 1 h incubation, the packed red cells were lysed by addition of three volumes of ice-cold water containing 1.4 mM neutral DTT as previously described [18]. Determination of rat liver microsomal and cytosolic GST activity was performed after appropriate dilution in 100 mM potassium phosphate buffer (pH 6.5). Since in the normal incubations with a final cytosolic protein concentration of $1.7 \text{ mg}\cdot\text{ml}^{-1}$, the cytosolic GST activity strongly exceeds the microsomal GST activity (approximately 95%), it was not necessary to remove the microsomes from these samples. When diluted cytosol with a final protein concentration of $0.2 \text{ mg}\cdot\text{ml}^{-1}$ was used the microsomes were sedimented from the cytosolic fraction by centrifugation at 4°C for 40 min at $110,000 \times g$. The clear supernatant was used for the determinations. The activity towards CDNB was determined as described by Habig and Jakoby [30]. The activities were measured at 25°C unless stated otherwise. Units of GST activity are defined as $1 \mu\text{mol}$ product produced per min.

Determination of GSH. For the determination of GSH in packed red cells and in rat liver cytosol after 1 h incubation, protein was precipitated by the addition of an equal volume of 8 % (w/v) trichloroacetic acid (TCA). To determine the GSH concentration in the erythrocyte incubations, hemoglobin was measured in the complete test system and in the packed red cells to evaluate the degree of concentration. Total GSH was determined in the TCA supernatants after dilution (9 fold) with 100 mM sodium/potassium phosphate buffer (pH 7.4) with the cyclic oxidation-reduction method essentially as described by Anderson [31].

Determination of methemoglobin formation. The percentage methemoglobin present in hemolysates was calculated from the absorbance change after addition of KCN compared to the same change in a sample fully converted to the methemoglobin form by the addition of $\text{K}_3\text{Fe}(\text{CN})_6$ [32].

Determination of lipid peroxidation products. The formation of lipid peroxidation products was assessed by determination of the amount of thiobarbituric-acid reactive substance present in the supernatant and was expressed as malondialdehyde (MDA) equivalents [33]. After incubation and centrifugation ($2000 \times g$), the supernatant was collected and protein was precipitated by the addition of an equal volume of 10% (w/v) TCA. 1.5 ml of the supernatant was added to 1 ml 1% (w/v) TBA solution in 5 mM NaOH and heated in a boiling water bath for 15 min. After forced cooling 1.25 ml butanol was added, and the two phases were thoroughly mixed. The absorbance of the butanol phase at 535 nm was determined and corrected for the background absorbance at 590 nm.

Statistical analyses

Significance of concentration dependent changes were analyzed separately for both metabolizing systems by linear regression using a model including day of experiment and drug concentration as explanatory variables. Using this model, we

tested whether slopes of parameter changes versus concentration were significant for each of the metabolizing systems, correcting for the day of the experiment. Effects were considered significant when the *p* value was smaller than 0.05 and when the effects were greater than 2.5% per mM. In microsomal incubations with CP and RS and in erythrocyte incubations with 3-HAA and MAS a clear saturation effect was found, in these cases no statistical analyses were performed. The regression model used, will correct for interexperiment variations in control values. Therefore, the standard errors (SE) shown in the figures are given as the standard error of the mean difference (SEMD) between sample value and its control (i.e. the corresponding value at concentration 0).

Difference between control values (i.e. without drug addition) of the two different metabolizing systems were analyzed using Wilcoxon's matched pair analysis with matches for control values from the same day.

The coefficients for day to day variation were calculated as $(SD/mean) \cdot 100\%$ for control GST activities and GSH concentrations. For the liver microsomal GST activity this was 4% of the control value ($0.125 \text{ U} \cdot \text{mg protein}^{-1}$), for liver cytosolic GST activity 10% (of $4.11 \text{ U} \cdot \text{mg protein}^{-1}$) and for erythrocyte GST activity 16.7% (of $5.20 \text{ U} \cdot \text{g Hb}^{-1}$). For the liver cytosolic GSH concentration this was 11.8% of the control value ($160.5 \mu\text{M}$) and for erythrocyte GSH concentration 15.7% (of $145.7 \mu\text{M}$).

Results

Microsomal incubations

Changes in rat liver microsomal GST activities in incubations with CP or c-DDP, with and without RS, are shown in figure 4.1. GST activities in equivalent incubations with MTX, 5-FU or Ara-C are given in table 4.1. Addition of RS alone (i.e. without drug addition) led to an increase of about 7% in microsomal GST activity ($p=0.0002$, $n=15$). For CP there was no significant change in GST activity in the absence of RS. In the presence of such a system the GST activity increased from $0.134 \text{ U} \cdot \text{mg protein}^{-1}$ (SE 0.001) without CP to $0.155 \text{ U} \cdot \text{mg protein}^{-1}$ (SE 0.004) at a CP concentration of 0.5 mM. Above this concentration no further increase in liver microsomal GST activity was seen for CP concentrations up to 2 mM. In c-DDP incubations a concentration dependent decrease in liver microsomal GST activity was found both with and without RS. The decreases amounted to $0.030 \text{ U} \cdot \text{mg protein}^{-1}$ (SEMD 0.002) and $0.018 \text{ U} \cdot \text{mg protein}^{-1}$ (SEMD 0.006) at 2.0 mM c-DDP with RS and without RS respectively. Significant GST decrease also occurred for MTX where the decreases amounted to $0.008 \text{ U} \cdot \text{mg protein}^{-1}$ per mM MTX and $0.011 \text{ U} \cdot \text{mg protein}^{-1}$ per mM MTX with RS and without RS respectively. Addition of 5-FU and Ara-C did not affect the liver microsomal GST activity, despite the presence of RS.

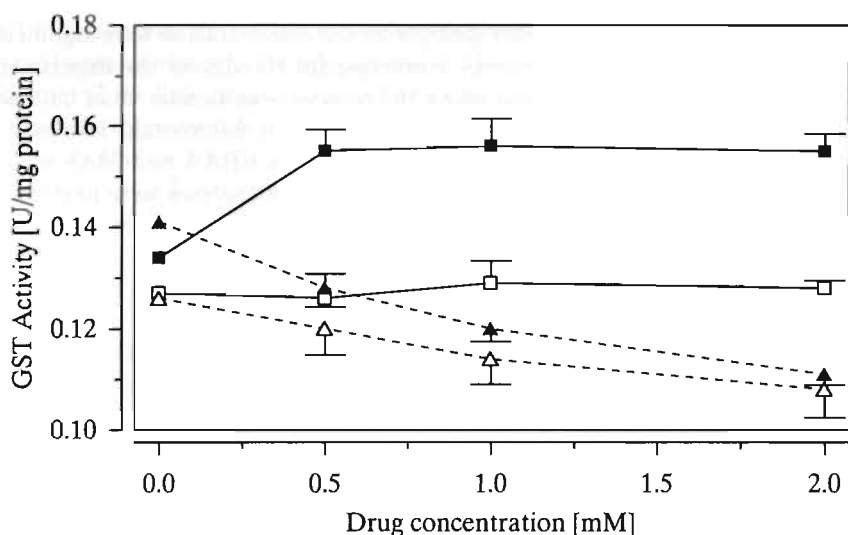


Figure 4.1: Effects of CP (\square) and c-DDP (\triangle) on the GST activity in rat liver microsomes after incubation (1 h, 37°C) with (closed symbols) and without (open symbols) a NADPH regenerating system (RS). The error bars represent the standard error of the mean difference (SEMD) between sample value and its control. GST inhibition was significantly related to the c-DDP concentration, both with RS ($p < 0.0001$) and without RS ($p = 0.0004$).

GST Activity (U-mg protein ⁻¹) in Rat Liver Microsomes			
Drug type	Drug concentration (mM)		
	0.0	1.0	2.0
MTX (RS) ^a	0.123	0.119 (0.004)	0.108 (0.006)
MTX ^a	0.112	0.095 (0.004)	0.090 (0.004)
5-FU (RS)	0.125	0.133 (0.004)	0.127 (0.004)
5-FU	0.120	0.125 (0.002)	0.121 (0.004)
Ara-C (RS)	0.131	0.124 (0.003)	0.130 (0.002)
Ara-C	0.124	0.127 (0.003)	0.121 (0.001)

a. GST inhibition was significantly related to the MTX concentration, both with RS ($p = 0.014$) and without RS ($p = 0.004$).

Table 4.1: Effects of MTX, 5-FU and Ara-C on the GST activity in rat liver microsomes after incubation (1 h, 37°C) with and without a NADPH regenerating system (RS). Values represent the mean followed by the standard error of the mean difference (SEMD) between sample value and its control in brackets.

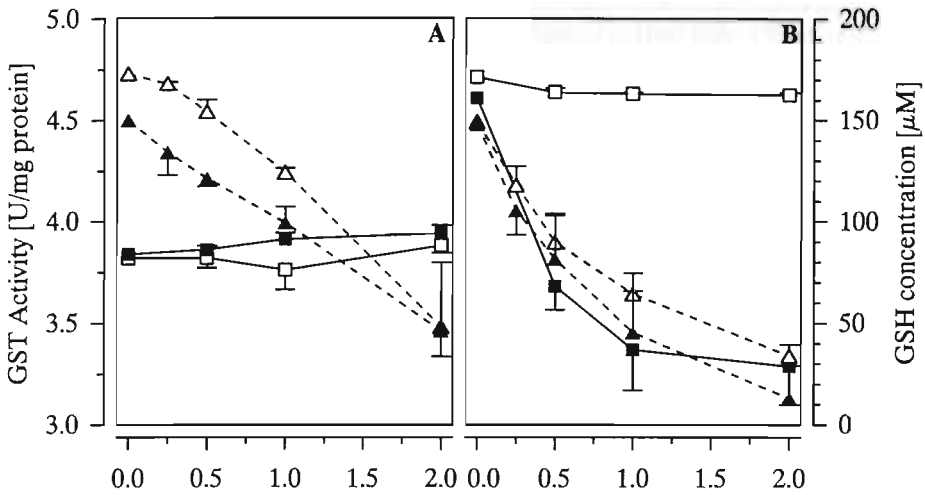


Figure 4.2: Effects of CP (\square) and c-DDP (\triangle) on the GST activity (A) and GSH concentration (B) in rat liver cytosol after incubation (1 h, 37°C) with (closed symbols) and without (open symbols) microsomal activating system (MAS). The error bars represent the standard error of the mean difference (SEMD) between sample value and its control. GST inhibition and GSH depletion were significantly related to the c-DDP concentration; $p < 0.0001$ for all the c-DDP effects. For CP only the GSH depletion in incubations with MAS was significantly related to the CP concentration; ($p = 0.004$)

Cytosolic incubations

Rat liver cytosolic GST (figure 4.2A) and GSH (figure 4.2B) values were determined in incubations with CP or c-DDP, with and without MAS. GST activities and GSH concentrations in equivalent incubations with MTX, 5-FU or Ara-C are given in table 4.2. In contrast to the increase seen in microsomal incubations after RS addition alone, here, addition of MAS alone (i.e. without drug addition) led to a decrease of about 9% in liver cytosolic GST activity ($p=0.0004$, $n=18$). Addition of MAS alone also led to a decrease of about 6% in GSH concentration ($p=0.005$, $n=12$). In c-DDP incubations a concentration dependent GST activity decrease was found of about $0.5 \text{ U} \cdot \text{mg protein}^{-1}$ per mM c-DDP both with and without MAS. Ara-C incubations with MAS resulted in a slight increase of GST activity ($0.12 \text{ U} \cdot \text{mg protein}^{-1}$ per mM Ara-C). In incubations without MAS Ara-C did not affect the GST activity. CP, MTX and 5-FU had no effect on the liver cytosolic GST activity despite the presence of MAS.

As CP and MTX were found to have an effect on the microsomal GST activity, we wondered whether a possible effect of CP and MTX on cytosolic GST might be prevented by the high GSH concentrations present. Therefore, incuba-

GST Activity (U·mg protein ⁻¹) in Rat Liver Cytosol			
Drug type	Drug concentration (mM)		
	0.0	1.0	2.0
MTX (MAS)	3.95	4.03 (0.14)	4.01 (0.06)
MTX	4.26	4.35 (0.01)	4.24 (0.06)
5-FU (MAS)	3.57	3.42 (0.10)	3.58 (0.15)
5-FU	3.68	3.85 (0.01)	3.75 (0.09)
Ara-C (MAS) ^a	3.95	4.07 (0.01)	4.19 (0.04)
Ara-C	4.26	4.32 (0.11)	4.42 (0.08)

GSH Concentration (μM) in Rat Liver Cytosol			
Drug type	Drug concentration (mM)		
	0.0	1.0	2.0
MTX (MAS)	132.3	133.9 (4.5)	135.6 (2.9)
MTX	146.8	142.9 (3.0)	146.1 (2.7)
5-FU (MAS)	163.3	151.9 (3.2)	162.7 (2.2)
5-FU	175.4	177.7 (15.2)	162.6 (12.2)
Ara-C (MAS)	132.3	126.9 (4.9)	130.2 (3.8)
Ara-C	146.8	143.9 (3.5)	141.2 (3.2)

a. GST inhibition in incubations with Ara-C and MAS was significantly related to the Ara-C concentration ($p = 0.0002$).

Table 4.2: Effects of MTX, 5-FU and Ara-C on the GST activity and GSH concentration in rat liver cytosol after incubation (1 h, 37°C) with and without microsomal activating system (MAS) Values represent the mean followed by the standard error of the mean difference (SEMD) between sample value and its control in brackets.

tions with diluted liver cytosol (final protein concentration 0.2 mg·ml⁻¹) were performed. In these incubations the remaining GSH concentration did not exceed the 15 μM level. The results of these incubations are given in table 4.3. Like what was found for the undiluted cytosol incubations, addition of MAS alone (i.e. without drug addition) had an inhibitory effect on the GST activity. This effect was stronger in the diluted cytosol incubations. A GST activity decrease of about 17% ($p=0.03$, $n=6$) was seen in the diluted cytosol incubations compared to a GST decrease of about 4% ($p=0.03$, $n=12$) in the concentrated cytosol incubations. The GST activity in incubations with CP and MAS actually increased from

GST Activity (U·mg protein ⁻¹) in Diluted Rat Liver Cytosol				
Drug type	Drug concentration (mM)			
	0.0	0.5	1.0	2.0
CP (MAS) ^a	4.56	5.56 (0.06)	5.63 (0.13)	5.72 (0.05)
CP	5.68	5.66 (0.04)	5.67 (0.13)	5.77 (0.14)
MTX (MAS)	3.59	— ^b	3.70 (0.08)	3.46 (0.02)
MTX	4.22	—	4.06 (0.04)	4.34 (0.14)

a. Increase of GST activity in incubations with CP and MAS was significantly related to the CP concentration ($p = 0.0007$).

b. Not determined

Table 4.3: Effects of CP and MTX on the GST activity in diluted rat liver cytosol (final protein concentration 0.20 mg·ml⁻¹) after incubation (1 h, 37°C) with and without microsomal activating system (MAS). GST activity in the CP incubates was measured at 37°C. Values represent the mean followed by the standard error of the mean difference (SEMD) between sample value and its control in brackets.

4.56 U·mg protein⁻¹ (SE 0.14) without CP to 5.72 U·mg protein⁻¹ (SE 0.12) at 2.0 mM CP and was linear over the whole range of concentrations tested. However, the activity value at 2.0 mM CP with MAS did not exceed the control value without MAS (5.68 U·mg protein⁻¹ (SE 0.14)). Without MAS there was no significant change in GST activity for CP. MTX had still no effect on the liver cytosolic GST activity despite the presence of MAS.

GSH was concentration dependently depleted in the non-diluted incubations by CP after metabolic activation and by c-DDP, both with and without MAS. In all these incubations, a GSH depletion of about 80% was found. CP incubations without MAS resulted in a minor GSH depletion (3.7 μ M per mM CP; $p=0.02$). Addition of 5-FU, MTX or Ara-C had no significant effect on the GSH concentration, despite the presence of MAS.

Erythrocytes incubations

Human erythrocyte GST (figure 4.3A) and GSH (figure 4.3B) values were determined in incubations with CP or c-DDP, with and without MAS. GST activities and GSH concentrations in equivalent incubations with MTX, 5FU or Ara-C are given in table 4.4. Similar to the liver cytosolic incubations, addition of MAS alone (i.e. without drug addition) led to a decrease of about 30% in erythrocyte GST activity and of about 19% in GSH concentration ($p=0.0001$ and $p=0.0002$ respectively, $n=15$). No significant change in GST activity was found in CP incubations without MAS. In the presence of MAS the GST activity actually increased from 4.31 U·g Hb⁻¹ (SE 0.58) without CP to 5.59 U·g Hb⁻¹ (SE 0.76) at 2.0 mM

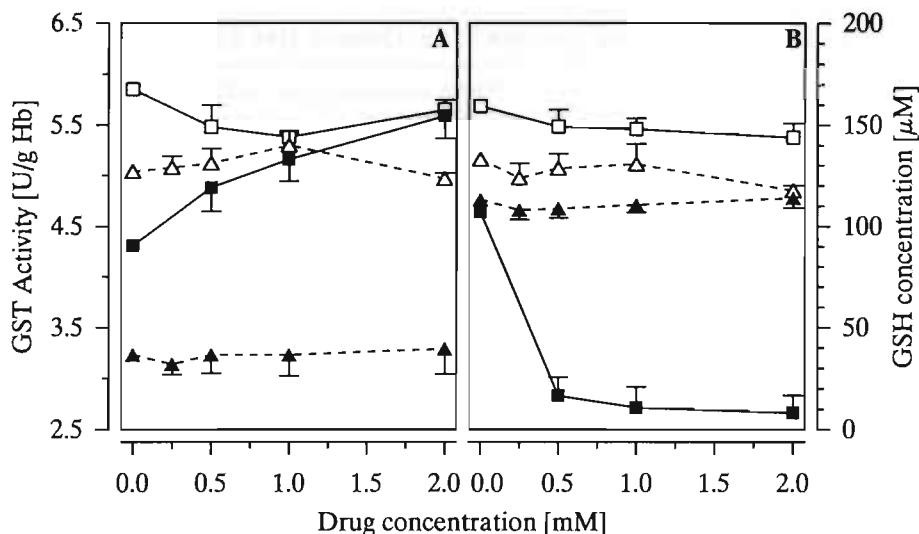


Figure 4.3: Effects of CP (\square) and c-DDP (\triangle) on the GST activity (A) and GSH concentration (B) in human erythrocytes after incubation (1 h, 37°C) with (closed symbols) and without (open symbols) microsomal activating system (MAS). The error bars represent the standard error of the mean difference (SEMD) between sample value and its control. Increase of GST activity and GSH depletion in incubations with CP and MAS were significantly related to the CP concentration ($p = 0.0004$ and $p = 0.02$ respectively).

CP and was linear over the whole range of concentrations tested. Here again, the activity value at 2.0 mM CP with MAS did not exceed the control value without MAS ($5.85 \text{ U} \cdot \text{g Hb}^{-1}$ (SE 0.59)). CP after metabolic activation depleted the GSH almost completely at a CP concentration of 0.5 mM (from $107.1 \mu\text{M}$ (SE 8.2) without CP to $16.6 \mu\text{M}$ (SE 2.8) at 0.5 mM CP). Although the decrease was clearly not linear, linear regression still showed a significant GSH depletion at increasing CP concentrations ($40.8 \mu\text{M}$ per mM CP). In CP incubations without MAS a small depletion in GSH was found ($6.8 \mu\text{M}$ per mM CP; $p=0.04$). In both types of incubations with c-DDP no effects were found on the GST activity and GSH concentrations. However, under the same conditions, GST inhibition values of 29% per mM c-DDP ($p=0.0001$) and 15% per mM c-DDP ($p=0.0001$) were observed in hemolysate and hepatocytes respectively. GSH was also concentration dependently depleted in hemolysate and hepatocytes (46% per mM c-DDP ($p=0.0001$) and 38% per mM c-DDP ($p=0.0001$) respectively). MTX, Ara-C and 5-FU had no effect on the erythrocyte GST activity and GSH concentrations despite the presence of MAS.

Since the increase in GST by addition of CP did not lead to values above the control value without MAS it is possible that the effects observed are not caused

GST Activity ($\text{U} \cdot \text{g Hb}^{-1}$) in Human Erythrocytes			
Drug type	Drug concentration (mM)		
	0.0	1.0	2.0
MTX (MAS)	3.61	3.50 (0.28)	3.37 (0.25)
MTX	4.89	4.82 (0.25)	4.89 (0.32)
5-FU (MAS)	3.42	3.18 (0.09)	3.22 (0.07)
5-FU	4.92	4.62 (0.14)	4.62 (0.18)
Ara-C (MAS)	3.61	3.62 (0.11)	3.53 (0.11)
Ara-C	4.89	4.89 (0.16)	4.89 (0.29)

GSH Concentration (μM) in Human Erythrocytes			
Drug type	Drug concentration (mM)		
	0.0	1.0	2.0
MTX (MAS)	120.3	109.0 (8.6)	109.9 (7.4)
MTX	143.4	140.3 (5.8)	148.9 (1.9)
5-FU (MAS)	113.4	111.5 (0.9)	115.8 (3.1)
5-FU	128.6	136.5 (4.9)	138.8 (11.0)
Ara-C (MAS)	120.3	110.9 (0.9)	109.2 (8.6)
Ara-C	143.4	138.8 (8.1)	132.9 (7.9)

Table 4.4: Effects of MTX, 5-FU and Ara-C on the GST activity and GSH concentration in human erythrocytes after incubation (1 h, 37°C) with and without microsomal activating system (MAS). Values represent the mean followed by the standard error of the mean difference (SEMD) between sample value and its control in brackets.

by real GST activation but rather by a decrease in the GST deactivation originating from MAS. To study this, three different experiments were performed. First, incubations with erythrocytes and higher CP concentrations were carried out. The purpose being to evaluate whether GST values above control values without MAS could be obtained in incubations with MAS and higher CP concentrations. Table 4.5 shows that this was not the case. Next we studied whether the effect of MAS could also be reduced with another substrate for microsomal activation. 3-Hydroxyacetanilide (3-HAA) was used for this purpose. 3-HAA is metabolized by MAS and the metabolites can deplete GSH in added erythrocytes [26]. No significant change in GST activity was found in 3-HAA incubations without MAS. Incubations with 3-HAA and MAS caused a GST activity *decrease* from $2.35 \text{ U} \cdot \text{g}$

GST Activity (U·g Hb ⁻¹) in Human Erythrocytes					
Drug type	Drug concentration (mM)				
	0.0	1.0	2.0	4.0	6.0
CP (MAS) ^a	2.15	— ^b	2.79 (0.03)	2.79 (0.04)	2.98 (0.10)
CP	3.36	—	3.17 (0.25)	3.07 (0.24)	3.14 (0.10)
3-HAA (MAS)	2.35	1.81 (0.23)	1.91 (0.10)	1.91 (0.14)	—
3-HAA	3.44	3.15 (0.18)	3.10 (0.06)	3.33 (0.13)	—

a. Increase of GST activity in incubations with CP and MAS was significantly related to the CP concentration ($p = 0.002$).

b. Not determined

Table 4.5: Effects of CP and 3-HAA on the GST activity in human erythrocytes after incubation (1 h, 37°C) with and without microsomal activating system (MAS). Values represent the mean followed by the standard error of the mean difference (SEMD) between sample value and its control in brackets.

Lipid Peroxidation and Methemoglobin Formation		
CP concentration (mM)	MDA release ^a (μ M)	Methemoglobin formation ^b (%)
0.0	0.11	0.05
0.5	0.11 (0.01)	-0.32 (0.29) ^c
1.0	0.11 (0.01)	-0.30 (0.17)
2.0	0.11 (0.01)	-0.37 (0.15)
0.0 (MAS)	29.56	3.48
0.5 (MAS)	29.08 (3.00)	4.36 (0.60)
1.0 (MAS)	23.93 (2.98)	4.74 (0.55)
2.0 (MAS)	23.53 (3.71)	4.52 (0.39)

a. Decrease in MDA release in incubations with MAS was significantly related to the CP concentration ($p = 0.01$).

b. Methemoglobin formation in incubations with MAS was significantly related to the CP concentration ($p = 0.03$).

c. The method used was specifically adapted to determine differences in methemoglobin formation. Absolute values are slightly higher and in all cases above zero.

Table 4.6: Effects of CP on MDA formation and methomoglobine formation in human erythrocytes after incubation (1 h, 37°C) with and without microsomal activating system (MAS). Values represent the mean followed by the standard error of the mean difference (SEMD) between sample value and its control in brackets.

Hb⁻¹ (SE 0.69) without 3-HAA to 1.81 U·g Hb⁻¹ (SE 0.46) at a 3-HAA concentration of 1.0 mM. No further decrease in human erythrocyte GST activity was seen for 3-HAA concentrations up to 4 mM (table 4.5). Finally, the effect of CP on lipid peroxidation (as indicated by MDA release) and formation of active oxygen species (AOS; as indicated by methemoglobin formation) were studied (table 4.6). MAS addition alone (i.e. without drug addition) clearly gave a large release of lipid peroxidation products (30 μ M SE 5.20) and gave methemoglobin formation (3.48 % SE 0.81; $p=0.03$, $n=6$ for both parameters). CP incubations without MAS showed no appreciable effect on MDA formation and methemoglobin formation. CP with MAS caused a minor concentration dependent decrease in MDA formation of 3.35 μ M per mM CP and a minor concentration dependent increase in methemoglobin formation of 0.45% per mM CP.

Discussion

Glutathione S-transferases and the non-protein thiol glutathione prevent the cell from damage through conjugation of electrophiles and removal of lipid peroxides. Various reactive chemicals influence the activity of the GSH/GST system, either by depletion of GSH or by modulation of GST activity [18,15,16,17]. Should such effects occur after *in vivo* drug treatment, this could lead to changes in the bioavailability of reactive compounds and by that to changes in drug efficacy. Furthermore, such changes in GSH content or GST activity could possibly be of use in biological effect monitoring.

The *in vitro* effects of the antineoplastic agents CP, c-DDP, MTX, 5-FU and Ara-C on the GSH/GST detoxification system were tested in rat liver fractions and in human erythrocytes. This *in vitro* test system allows to discriminate whether compounds or the metabolites formed are able to enter the erythrocyte [26]. Of the five antineoplastic agents used in this *in vitro* study, CP and c-DDP are known to act as alkylating compounds, while 5-FU, Ara-C and MTX are known to act as antimetabolites. Therefore, effects of CP and c-DDP on the GSH/GST system were more likely *a priori* than effects by 5-FU, Ara-C and MTX.

Addition of RS alone (i.e. without drug addition) led to an increase in liver microsomal GST activity. In contrast, addition of MAS alone led to a decrease in liver cytosolic and in erythrocyte GST activity and GSH concentrations. This effect in the control values is probably a result of AOS formed by the microsomes in the presence of RS [34]. Supportive evidence for increased AOS availability in erythrocytes was given by methemoglobin formation and lipid peroxidation in erythrocyte incubations with MAS (table 4.6). It is known that microsomal GST activity and the soluble class μ GST can be activated by AOS [35,36,37,38]. GSTs from the α and π class can be inactivated by AOS [38,39,40,41]. Erythrocytes only contain the π class GST, while the α class GSTs are the most abundant

in rat liver cytosol, especially when rats are pretreated with phenobarbital [2]. From this one can conclude that AOS generated by MAS probably caused the initial increases in microsomal GST activity and the initial decreases in rat liver cytosolic GST (α) activity and in human erythrocyte GST (π) activity. The initial GST decreases in incubations with non-diluted cytosol were much smaller than in incubations with diluted cytosol. This suggests that GSH prevent GST activity changes by AOS. Depletion of GSH by AOS in the control incubations with MAS may be due to direct oxidation, and to consumption in protective reactions catalyzed by GST and glutathione peroxidase [12,13]. Furthermore, the liver and erythrocytes export oxidized glutathione and GSH S-conjugates [12,13] and oxidation of GSH leads to mixed disulfide formation with proteins [42].

5-FU did not affect the GST activity and GSH concentrations in the different incubations. For Ara-C with MAS a slight increase in cytosolic GST activity was found. As other incubations with Ara-C expressed no clear effect, and since it is known that Ara-C requires no microsomal metabolism, it is likely that Ara-C does not affect GSH or GST levels and that the single divergent result was found by chance. The microsomal GST enzyme is different from soluble GST enzymes with regard to occurrence, substrate specificity, kinetic parameters [43] and in fact belongs to a different gene family [44]. Our results (table 4.1) indicate that MTX at millimolar concentrations is only able to inhibit about 20% of the microsomal GST activity. As expected in advance no serious effects were found for the three antimetabolites.

In vivo reduction of GST activity in liver and kidney was previously described for c-DDP [45]. Next to this, c-DDP reacts with sulfhydryl groups of enzymes and sulfur containing amino acids, peptides and proteins such as cysteine, GSH and metallothionein [29,46,47,48]. This is consistent with our findings where c-DDP causes a concentration dependent GST inactivation and GSH depletion in cytosolic liver incubations. However, in erythrocyte incubations there was no significant effect of c-DDP on GSH or GST levels. This implicates that c-DDP is not able to enter the erythrocyte under the conditions used. This was not due to the use of PBS in stead of sodium/phosphate buffer, as incubations with erythrocytes and this buffer also gave negative results. In fact PBS was used because the two chloride ligands are stable at the chloride concentration of the extracellular matrix. After diffusion into a cell, the lower chloride concentration permits loss of chloride from the drug. The drug becomes then an aquated, charged electrophile which reacts with nucleophilic sites on cellular macromolecules [29]. Moreover, c-DDP caused a decrease of GSH and GST levels in hemolysate and in hepatocytes. A possible explanation for the phenomenon that c-DDP does hardly pass the erythrocyte membrane is the fact that c-DDP binds avidly to red cell membranes [49,50].

CP is metabolized mainly by hepatic microsomal enzymes to 4-hydroxycyclophosphamide, producing equimolar amounts of phosphoramidate mustard and acrolein [51]. CP metabolites, especially acrolein, have long been known to cause

GSH depletion [52,53]. As expected GST activities were not affected in any of the CP incubations without MAS. However, CP incubations without MAS induced a slight GSH depletion in rat liver cytosol and in human erythrocytes. This indicates that erythrocytes and liver cytosol itself may be able to activate a small amount of CP. GSH was almost completely depleted and GST activity was increased in all incubations with CP in the presence of MAS. This confirms that CP has to be metabolized to give an effect and proves that the CP metabolites can enter the erythrocyte. Activation of microsomal GST was already at a maximum at a CP concentration of 0.5 mM. It is known that acrolein activates microsomal GST if there is no GSH available [54], and since acrolein is a major CP metabolite [22,51], the microsomal GST activation seen here was probably caused by acrolein. No increase in GST activity was seen after CP treatment with MAS in the liver cytosolic incubations. After dilution of the cytosol, a clear GST increase was found, suggesting that GSH prevent GST activity changes by CP metabolites. Moreover, when GSH was added to incubations with diluted cytosol the GST activation was lowered (data not shown). The increase in GST activity seen in diluted liver cytosol and erythrocyte incubations after CP treatment with MAS did not exceed the control value without MAS. Therefore, it could not be excluded that this increase was actually a reduction of the inhibition caused by MAS addition. So, perhaps CP serving as a substrate for the microsomes, caused a reduction in AOS formation leading to a reduction in GST inactivation. To detect whether the GST activity changes were caused by reactive CP metabolites or by a reduction in AOS, three additional experiments were performed. As a first approach, higher concentrations CP were used (up to 6 mM). However, even under these conditions no GST activities above the control value were found. 3-HAA, another substrate for the microsomes, showed no concentration dependent increase in GST activity. In other words there was no significant reduction in the formation of AOS responsible for GST inactivation. Therefore, reduction in AOS formation by microsomal substrates does not seem to be a general phenomenon. Finally, methemoglobin formation and lipid peroxidation were evaluated. The rational being that CP induced reduction in AOS formation should lead to concomitant changes in these parameters. A CP concentration dependent decrease in MDA release was found in the incubations with MAS. However, the relative change in MDA release is much smaller than the relative change in GST activity. Contrary to what might be expected if CP caused a reduction in AOS formation, CP caused a concentration dependent *increase* in methemoglobin formation. Based on the latter three findings it is not likely that a reduction in AOS formation is responsible for the activation of GST but rather that the CP metabolites are responsible for the GST activation.

The GST activity and the GSH concentrations in tumor cells are thought to be very important for the resistance against chemotherapy [5,6,7,8,9] and radiation therapy [13,14]. GSH and GST detoxify some alkylating antineoplastic agents and detoxify organic hydroperoxides formed as a result of radical for-

mation during treatment with some antineoplastic agents and during radiation therapy [1, 4]. Therefore, changes in GST and GSH levels may result in different cytotoxicity patterns and may influence drug efficacy. The results presented here indicate that apart from changes in GST expression, direct modulation of the available enzyme may be important. Since modulation will be isoenzyme specific and the GST isoenzyme content in tumor cells is known to differ from normal tissue [8], this modulation, particularly seen here in erythrocytes, is likely to play a role in relative sensitivity of tumor tissue. In addition, tumors could be sensitized to alkylating antineoplastic agents and radiation therapy by GSH depletion. In fact, GSH depletion with L-buthionine sulfoximine and diethylmaleate has been suggested for this purpose [55, 56]. So, GSH depletion in tumors by alkylating antineoplastic agents in combination with radiation therapy will be a useful strategy for tumor sensitization. This would especially be the case if targetting of GSH depleting and/or GST inactivating antineoplastic agents could result in selective sensitization of tumor tissue. The two alkylating antineoplastic agents used in this study both resulted in significantly lowered GSH levels and c-DDP also caused GST activity decrease. CP on the other hand increased the GST activity but this increase is negligible in comparison with the GSH depletion. Therefore, changes in GST activity and in GSH levels induced by CP and c-DDP may result in increased tumor sensitivity to subsequent radiation therapy.

Acknowledgements The authors wish to thank Vivian Geelen and Raphael Janssen for their technical support, Dr A. Kester for his advice in statistical analyses, the Red Cross Blood Bank "Zuid Limburg" for the collection of blood samples and Multipharma B.V. (Weesp, NL) for their generous gifts (cis-diamminedichloroplatinum(II) and methotrexate).

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Chapter 5

Two Mechanisms for Toxic Effects of Hydroxylamines in Human Erythrocytes: Involvement of Free Radicals and Risk of Potentiation

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Summary. The toxic potency of three industrially used hydroxylamines was studied in human blood cells *in vitro*. The parent compound hydroxylamine (HYAM) and the O-ethyl derivative (OEH) gave very similar results. Both compounds induced a high degree of methemoglobin (Hb^+) formation and glutathione depletion. Cytotoxicity was visible as Heinz body formation and hemolysis. High levels of lipid peroxidation (LP) occurred, in this respect OEH was more active than HYAM. In contrast H_2O_2 induced LP was lowered after OEH or HYAM treatment, this is explained by the ferrohemoglobin dependence of H_2O_2 induced radical species formation. Glutathione S-transferase (GST) and NADPH methemoglobin reductase (NADPH-HbR) activities were also impaired, probably as a result of the radical stress occurring. The riboflavin availability was decreased. Other enzyme activities (glutathione reductase (GR), glucose 6-phosphate dehydrogenase (G6PDH), glucose phosphate isomerase and NADH methemoglobin reductase) were not or only slightly impaired by HYAM or OEH treatment.

A strikingly different scheme of reactivity was found for N,O-dimethyl hydroxylamine (NODMH). This compound gave much less Hb^+ formation and no hemolysis or Heinz body formation at concentrations up to and including 7 mM. LP induction was not detectable, but could be induced by subsequent H_2O_2 treatment. GST and NADPH-HbR activities and riboflavin availability were not decreased. On the other hand GR and G6PDH activities were inhibited. These results combined with literature data indicate the existence of two different routes of hematotoxicity induced by hydroxylamines. HYAM as well as O-alkylated derivatives primarily induce Hb^+ , a process involving radical formation. The radical stress occurring is probably responsible for most other effects. N-alkylated species like NODMH primarily lead to inhibition of the protective enzymes G6PDH and GR. Since these enzymes play a key role in the protection of erythrocytes against oxidative stress a risk of potentiation during mixed exposure does exist.

Introduction

Hydroxylamines are derivatives of the parent compound hydroxylamine (HYAM, chemical formula H_2NOH). Hydroxylamine itself is used in the synthesis of Caprolactam, which is used for the production of Nylon 6TM. It is also used as a reducing agent in photography, as an antioxidant for soaps and fatty acids, as a tanning agent and as an intermediate in the production of pesticides and pharmaceuticals [8, 30]. Derivatives of HYAM often serve as intermediates in chemical synthesis. The main toxic effect of HYAM is of hematological nature [8, 30]. Methemoglobin (Hb^+) formation was found *in vitro* [7, 11, 18, 63]. In *in vivo* animal studies anemia with increased Hb^+ levels and sometimes with sulfhemoglobinemia was consistently found [3, 8, 20, 30, 39, 44, 54, 59, 60, 71, 72]. In some studies this was

found to be accompanied by remarkable splenomegaly [20, 54, 72]. In a more recent subchronic (ninety days) study —performed according to guidelines of the Organisation for Economic Co-Operation and Development (OECD)— rats were exposed to HYAM-sulfate [5]. In the 250 ppm exposed groups (male and female) hematological effects and increased spleen weight were found among other effects. In the 50 ppm treated groups small increases in reticulocyte numbers and some other marginal effects were found. The “no effect level” in this study was between 10 and 50 ppm for both sexes. In his review Gross [30] quotes a single incidence of human hydroxylamine poisoning described in Russian literature [45]. A woman who had drunk about “two swallows” of HYAM solution in water developed rapidly progressing hemolytic anemia which required five weeks for recovery.

Martin *et al.* [46] described the development of serious anemia in five laboratory workers working with methylated hydroxylamines (O-methyl, N,O-dimethyl (NODMH) and trimethyl hydroxylamine). In the most acute patient lowered hemoglobin and increased serum iron levels were found, indicating hemolytic anemia. All patients had shown jaundice symptoms and had increased numbers of bone marrow erythroblasts. Reticulocyte numbers were increased in two patients. Heinz bodies were not found and Hb⁺ could not be detected spectrophotometrically. These findings prompted the authors to further *in vitro* studies.

Interestingly trimethyl hydroxylamine in these studies gave by far the highest hemolytic activity, while it did not lead to increased methemoglobin levels. In contrast the far less hemolytic O-methyl hydroxylamine and the N,O-dimethyl compound did strongly induce methemoglobin formation. The higher hemolytic potency of trimethyl, and to a much lesser extent N,O-dimethyl hydroxylamine, was found to be accompanied with strong glucose 6-phosphate dehydrogenase (G6PDH) inhibition and loss of reduced glutathione (GSH).

This suggests that two separate mechanisms for the hematotoxicity of hydroxylamines may exist. For the first one methemoglobin formation is a major effect, for the other G6PDH inhibition is predominant. Both routes seem to be able to give rise to hemolytic anemia. The purpose of the present study was to further elucidate these two mechanisms. Three industrially used hydroxylamines were tested: HYAM, O-ethyl hydroxylamine (OEH) and NODMH. Next to the effects described above we were especially interested in the fact that the Hb⁺ formation by HYAM does involve a radical mechanism [63], and therefore might give rise to free radical induced lipid peroxidation.

Materials and Methods

Chemicals and Enzymes

The following chemicals were used: N,O-dimethyl hydroxylamine (NODMH, methoxymethylamine, CAS-nr 1117-97-1, purity 99.5%) and O-ethyl hydroxylamine (OEH, ethoxyamine, CAS-nr 624-86-2, 55% (w/v) solution in water),

were a gift from DSM Special Products (Geleen, NL). Brilliant cresyl blue was from BDH (Poole, UK), malonaldehyde bis(diethylacetal) (used as precursor for malondialdehyde reference) was from Janssen Chimica (Beerse, Belgium), 2-thiobarbituric-acid (*TBA*) was from Merck (Darmstadt, FRG), 1-chloro-2,4-dinitrobenzene, 5,5'-dithio-bis(2-nitrobenzoic acid), 4,4'-dithiodipyridine (*4-PDS*), dithiothreitol, flavin adenine dinucleotide (*FAD*), fructose 6-phosphate, glucose 6-phosphate, glucose 6-phosphate dehydrogenase (grade XI from *Torula* yeast) (*G6PDH*), reduced (*GSH*) and oxidized (*GSSG*) glutathione, glutathione reductase (type III) (*GR*), hydroxylamine HCl (*HYAM*, CAS-nr: 5470-11-1 (free base CAS-nr: 7803-49-8) (ACS reagent, purity 99.1%), methylene blue, reduced β -nicotinamide adenine dinucleotide (*NADH*), β -nicotinamide adenine dinucleotide phosphate (*NADP*), reduced β -nicotinamide adenine dinucleotide phosphate (*NADPH*), and 6-phosphogluconic dehydrogenase were from Sigma (St. Louis, Mo, USA). All other chemicals were of analytical quality. Only microfiltered deionized water was used.

Blood samples.

Human blood samples were collected in vacuum tubes containing K_3EDTA as an anticoagulant. For the determinations of Heinz body formation, lipid peroxidation, peroxidation resistance and glutathione S-transferase and glutathione reductase activities fresh samples from individual persons were used. For all other experiments pooled blood samples from three to five persons with blood group O+ were used. The latter samples were checked for viral infections and stored at 4°C until the next day.

Incubations

To 1 ml blood 100 μ l hydroxylamine solution in 100 mM KH_2PO_4/Na_2HPO_4 of pH 7.4 were added, and the samples were incubated in a shaking waterbath (80 revolutions per minute) for 1 hour at 37°C. All experiments were performed three times on separate days and with different blood samples, each day all incubations were performed in duplicate. Except for the determinations of Heinz bodies, hemolysis and lipid peroxidation, erythrocytes were washed three times with five volumes of cold phosphate buffered saline (15 mM KH_2PO_4/Na_2HPO_4 + 130 mM NaCl; pH=7.4). Where applicable the erythrocytes were lysed by addition of three to five volumes of cold water, after 15 minutes storage on ice cellular debris was removed by centrifugation at $13000\times g$ for 10 min.

Analyses

Heinz body formation. To 1 ml blood 0.5 ml 1% (w/v) brilliant cresyl blue in saline was added. Heinz bodies are colored during subsequent incubation. More severe cellular damage leads to detectable numbers of Heinz bodies after shorter incubation times and at lower temperatures. For this reason, of two identical

samples one was incubated at 20°C and the other at 37°C. After 20, 40 and 60 minutes of incubation samples were dried and evaluated by light microscopy (1000×) under oil immersion [33].

Hemolysis. After incubation the Hb content of the plasma was used to determine the degree of hemolysis.

Methemoglobin (Hb^+). The percentage Hb^+ present in hemolysates was calculated from the absorbance change after addition of KCN compared to the same change in a sample fully converted to the Hb^+ form by the addition of $\text{K}_3\text{Fe}(\text{CN})_6$ [65].

Total glutathione (GT). GT was determined after precipitation of protein (200 μl packed cells plus 200 μl 8% (w/v) trichloroacetic acid) using the cyclic oxidation-reduction method essentially as described by Anderson [2]. Especially for the determination of GT, it was of interest whether the *Torula* yeast GR enzyme used was itself inhibited by any of the hydroxylamines studied. This was tested separately with hydroxylamine concentrations of 0, 2.5 and 7 mM, no significant inhibition of *Torula* yeast GR was found under the assay conditions used.

Reduced glutathione (GSH). GSH was determined as non-protein sulfhydryl after trichloroacetic acid precipitation of erythrocytes by measurement of the reactivity towards 5,5'-dithio-bis(2-nitrobenzoic acid) as previously described [23].

Release of thiobarbituric-acid reactive substances (TBARS). The accumulation of lipid peroxidation (LP) products in the extracellular medium was assessed by determination of the amount of TBARS released and was expressed as malondialdehyde equivalents [62]. After incubation of 1.5 ml fresh blood with hydroxylamines, the plasma was collected after centrifugation (1500 \times g for 10 min). The protein was precipitated by addition of an equal volume of 15% (w/v) trichloroacetic acid and subsequent centrifugation. 1.5 ml of the supernatant was added to 1 ml TBA solution (1% (w/v) in 50 mM NaOH) and heated in a boiling waterbath for 15 min. After forced cooling, 1.25 ml butanol was added, and the two phases were thoroughly mixed. The absorbance of the butanol phase at 535 nm was determined and corrected for the background absorbance at 590 nm.

Resistance to H_2O_2 forced lipid peroxidation. The remaining resistance to H_2O_2 induced LP of erythrocytes pretreated with hydroxylamines was assessed. After incubation of fresh blood samples with hydroxylamines the plasma was removed by centrifugation. The erythrocytes were washed three times to remove remaining hydroxylamines and were diluted in phosphate buffered saline to a cell concentration of 5% (v/v). To 2.5 ml erythrocyte suspension an equal volume of 20 mM

H₂O₂ plus 1 mM NaN₃ was added. TBARS release was determined after reincubation at 37°C for 0, 15, 30, 45 and 60 min [22].

Free hemoglobin sulfhydryl (HbSH). Free hemoglobin sulfhydryl groups were determined in hemolysates from the reactivity towards 4-PDS [29] as previously described [23].

NADH methemoglobin reductase (NADH-HbR). NADH-HbR activity in hemolysates was determined using DEAE-cellulose purified hemoglobin converted to the methemoglobin-ferrocyanide complex as substrate with the method described by Hegesh *et al.* [35] and some small modifications according to Bauer [6].

NADPH methemoglobin reductase (NADPH-HbR). NADPH-HbR activity in hemolysates was determined from the reduction of methylene blue by NADPH [9,38]. Disappearance of NADPH was followed spectrophotometrically at 340 nm. Corrections were made for spontaneous methylene blue reduction.

Glutathione S-transferase (GST). GST (EC 2.5.1.18) activity with 1-chloro-2,4-dinitrobenzene as substrate was determined in hemolysates prepared from incubates of fresh blood samples by the addition of three volumes 1.4 mM neutralized dithiothreitol, using the method of Habig and Jakoby [32] with previously described modifications [10].

Glutathione reductase (GR). GR (EC 1.6.4.2) activity in hemolysates was determined by a modification of the method described by Carlberg and Mannervik [12]. 100 μ l Hemolysate were added to 2 ml 100 mM potassium phosphate buffer, 50 μ l 80 mM EDTA, 100 μ l 2 mM NADPH and 100 μ l 0.3 mM FAD. After 2 min pre-incubation (37°C), the reaction was started by addition of 100 μ l 7.5 mM GSSG. After 15 sec the reaction was followed spectrophotometrically at 340 nm for 2 min. The FAD was added in order to convert all enzyme to its holo form.

Glutathione reductase riboflavin activity coefficient (GR_{coeff}). The coefficient of erythrocyte GR activity with and without addition of FAD to the assay was used as an indicator of the riboflavin content. The determinations were carried out exactly as described for GR above except that FAD was left out from one of the 2 samples[65].

Glucose 6-phosphate dehydrogenase (G6PDH). G6PDH (EC 1.1.1.49) activity in hemolysates was determined spectrophotometrically; the reduction of NADP⁺ was followed at 340 nm [65].

Glucose-phosphate isomerase. Glucose-phosphate isomerase (EC 5.3.1.9) activity in hemolysates was determined using fructose 6-phosphate as substrate in the presence of NADP⁺, G6PDH and 6-phosphogluconic dehydrogenase. Under these circumstances each molecule of fructose 6-phosphate converted will give rise to reduction of two molecules NADP⁺, which was followed spectrophotometrically at 340 nm [6].

Hemoglobin (Hb). The Hb concentrations were determined with the hemoglobin cyanide procedure [41].

Statistical Evaluation

Significance of concentration dependent changes was evaluated using linear regression analysis with a model including day of experiment and hydroxylamine concentration as explanatory variables. Using this model, we tested whether slopes of parameter changes versus concentration were significant correcting for the day of experiment. In each experiment incubations were performed with 0, 1, 2.5, 5 and 7 mM hydroxylamine concentrations. All experiments were performed three times. Duplicate values obtained during a single experiment were averaged before statistical analysis since these values are not independent. This leads to a total of 3 (experiments) \times 5 (concentrations) = 15 data points for each regression analysis. The regression model used, will correct for interexperiment variations in control values. For this reason the standard errors shown in the figures are given as the standard error of the mean difference between sample value and its control (i.e. the corresponding value at concentration 0).

Results

Oxidative effects

A concentration dependent Hb⁺ formation was found for all three hydroxylamines (Figure 5.1 on the following page). The effects of HYAM and OEH (about 8% Hb⁺ formation for each mM addition) were much stronger than the effect caused by NODMH. GT decreased for all three hydroxylamines (Figure 5.3A on page 85). GT depletion caused by OEH (45 (SE 2.5) μ M GT decrease/mM OEH) was much more pronounced than the effect caused by the other two hydroxylamines (24 (SE 3) and 19 (SE 3) μ M GT decrease/mM for HYAM and NODMH respectively). The decreases in reduced GSH also were strongest for OEH (Figure 5.3B). The regression coefficient of this GSH decrease amounted to 36 (SE 3) μ M/mM OEH and to 18 (SE 3) μ M/mM HYAM. For NODMH the depletion was only about 8 (SE 4) μ M/mM, and the significance of this depletion was doubtful ($p = 0.0726$). Plasma GT was found to increase for all three hydroxylamines (figure 5.3C). Contrary to what might be expected from erythrocyte depletions, the increase

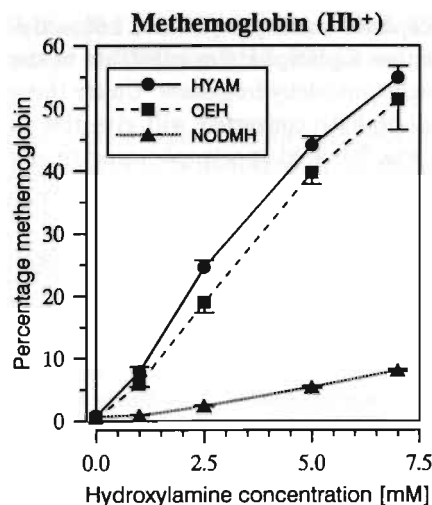


Figure 5.1: Formation of methemoglobin in human erythrocytes after incubation with hydroxylamines (1 hour, 37°C). Regression analyses showed significant ($p < 0.0001$) increases in methemoglobin formation with increasing concentration for all three hydroxylamines.

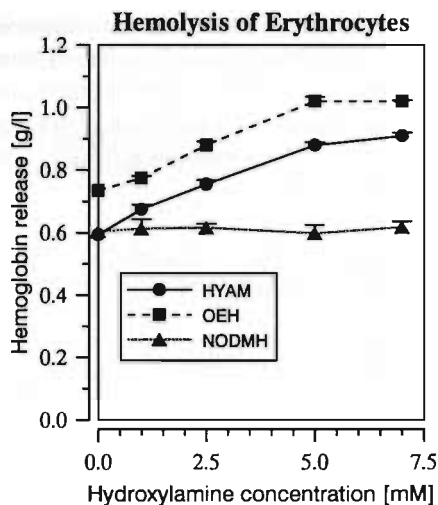


Figure 5.2: Occurrence of hemolysis of human erythrocytes after incubation with hydroxylamines (1 hour, 37°C). Regression analyses showed significant ($p < 0.0001$) increases in hemolysis with increasing HYAM and OEH concentrations.

was strongest for NODMH (about 5.4 μM increase/mM NODMH). For OEH and HYAM increases were not more than half that amount. Loss of hemoglobin sulfhydryl group availability was not found for any of the chemicals studied.

Cell damage

Increased hemolysis occurred for HYAM and OEH (Figure 5.2). Both compounds liberated about 45 mg Hb/l from the erythrocytes for each mM added. For NODMH no increased hemolysis was detectable. Heinz bodies were visible in all blood samples treated with 2.5 mM HYAM or OEH (Table 5.1 on the next page). In the 2.5 mM HYAM treated samples cell blue spots, normally indicating Heinz bodies, were also visible extracellularly. The same indications for cell breakdown were also present in the 2.5 mM OEH treated samples, but only when they were incubated at 37°C. Minor Heinz body formation was present in the 1 mM HYAM treated samples incubated at 37°C or incubated for 60 min or more at 21°C as well as in the 1 mM OEH treated samples incubated at 37°C for 60 min or more. The occurrence of Heinz bodies at a lower temperature and after shorter incubation times for HYAM compared to OEH, indicates that the effect is strongest for HYAM. In NODMH treated blood samples no Heinz body formation could be detected even at concentrations up to 10 mM.

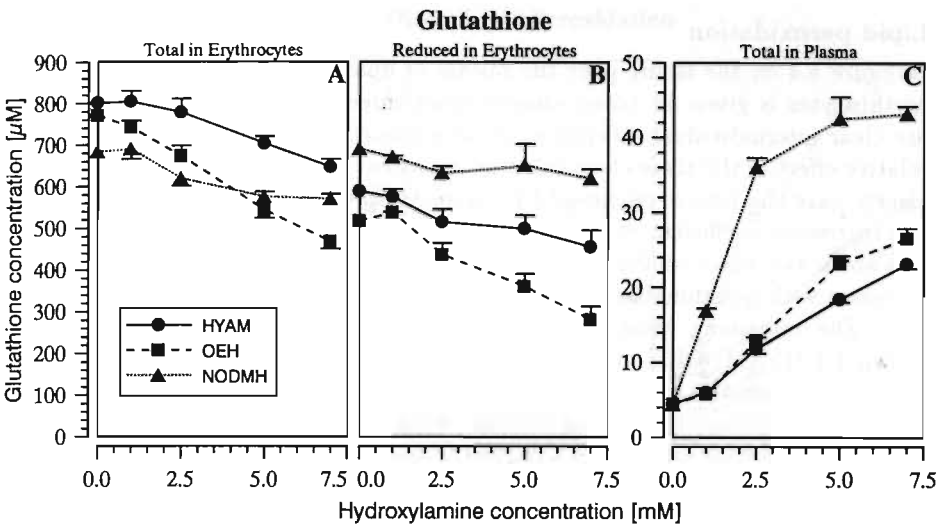


Figure 5.3: Availability of A: total glutathione (reduced plus oxidized) in human erythrocytes; B: reduced glutathione in human erythrocytes and C: total glutathione in human plasma, after incubation with hydroxylamines (1 hour, 37°C). Separate experiments were performed to obtain the data for panel A, B and C. Regression analyses showed that all concentration dependent changes were significant with $p < 0.0001$, with the exception of the decrease in reduced glutathione with increasing NODMH (panel B) where $p = 0.073$.

Sample	21°C			37°C		
	20 min	60 min	120 min	20 min	60 min	120 min
Blank	-	-	-	-	-	-
NODMH 10 mM	-	-	-	-	-	-
HYAM 1 mM	-	±	±	±	±	+
HYAM 2.5 mM	+ ¹	+ ¹	+ ¹	+ ¹	+ ¹	+ ¹
OEH 1 mM	-	-	-	-	±	±
OEH 2.5 mM	+	+	+	+ ¹	+ ¹	+ ¹

¹ Blue spots were also seen extracellularly.

Table 5.1: Presence of Heinz bodies in human erythrocytes after incubation with hydroxylamines (1 hour, 37°C). After addition of brilliant cresyl blue the erythrocytes were incubated at 21°C or 37°C for a specified amount of time. Visibility of Heinz bodies is indicated as positive (+), negative (-) or visible but minor (±).

Lipid peroxidation

In figure 5.4 on the facing page the release of lipid peroxidation products from erythrocytes is given for blood samples from three individuals. Although there are clear interindividual differences in both susceptibility and background, the relative effect of the three chemicals was consistent in all three individuals. OEH clearly gave the largest release of LP products. For the three persons combined the regression coefficient was 117 (SE 13) nM/mM OEH. The effect of HYAM was about two times smaller, but still very significant. NODMH gave only minor increases with questionable statistical significance ($p = 0.093$).

The remaining resistance of erythrocytes towards H_2O_2 induced LP is shown for HYAM, OEH and NODMH in figures 5.5, 5.6 and 5.7 respectively. Fresh blood samples from three individuals were used, and the individual data are presented separately. As can be seen clearly from the figures the main effect of hydroxylamine treatment is a changed speed of TBARS liberation (i.e. the slopes of the TBARS release versus time curves change). The patterns seen for HYAM (figure 5.5) and OEH (figure 5.6) are very similar. For both substances and in the samples from all three individuals pretreatment with one of these two hydroxylamines leads to a hydroxylamine-concentration dependent decrease in the H_2O_2 induced LP. For NODMH the effect occurring is more complex. The amounts of LP products released from erythrocytes from person A were not markedly influenced by NODMH pretreatment. Blood samples from this individual did generally have the lowest induced LP both after direct treatment with hydroxylamines (figure 5.4) and after subsequent H_2O_2 treatment (figures 5.5 and 5.6). The erythrocytes from the other two individuals showed a hydroxylamine-concentration dependent increase in H_2O_2 induced LP after 1 and 2.5 mM pretreatment. In samples from person B however, the response to H_2O_2 was lower in the 5 and 7 mM pretreated samples than the (very high) response in the 2.5 mM pretreated samples.

Enzyme inhibitions

GST activity was strongly decreased in both HYAM and OEH treated samples (figure 5.8 on page 89). For HYAM and OEH the GST activity decreased with about $0.35 \text{ U} \cdot \text{g Hb}^{-1} / \text{mM}$ addition. As a consequence about two-thirds of the initial GST activity was lost at the highest ($=7 \text{ mM}$) HYAM and OEH concentrations used. For NODMH no significant changes in the GST activity were found. In contrast the GR activity was only decreased for NODMH, for this substance GR activities in 7 mM incubations were about 20% lower than the $6.7 \text{ U} \cdot \text{g Hb}^{-1}$ found in control incubations (figure 5.10A on page 90). The GR_{coeff} was slightly decreased in both HYAM ($p=0.025$) and OEH ($p=0.01$) treated samples (figure 5.10B). This indicates a decrease in riboflavin availability in these samples [65].

The G6PDH activity was strongly impaired by NODMH ($0.76 \text{ (SE } 0.04) \text{ U} \cdot \text{g}$

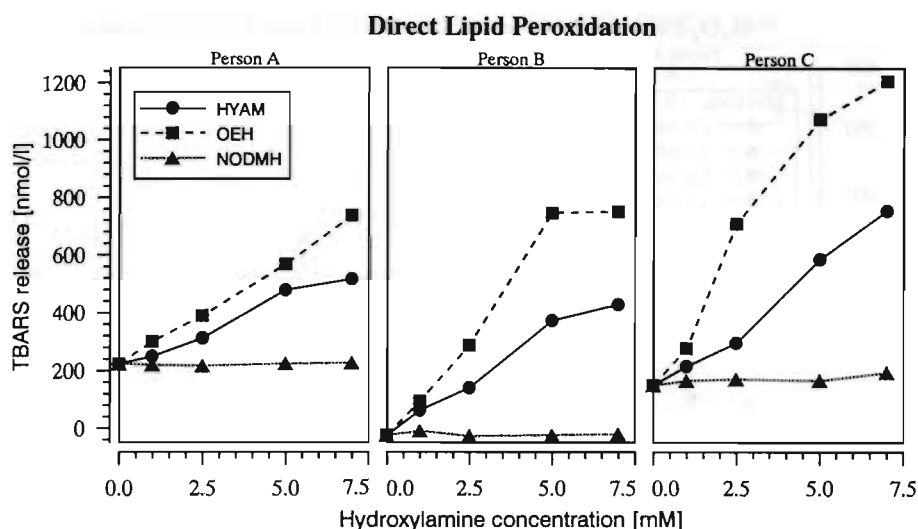


Figure 5.4: Lipid peroxidation, expressed as malondialdehyde equivalents in reactivity towards TBA, in blood samples of three individuals after incubation with hydroxylamines (1 hour, 37°C). Regression analyses on the combined data showed significant increases in TBARS release with increasing OEH and HYAM concentrations ($p < 0.0001$).

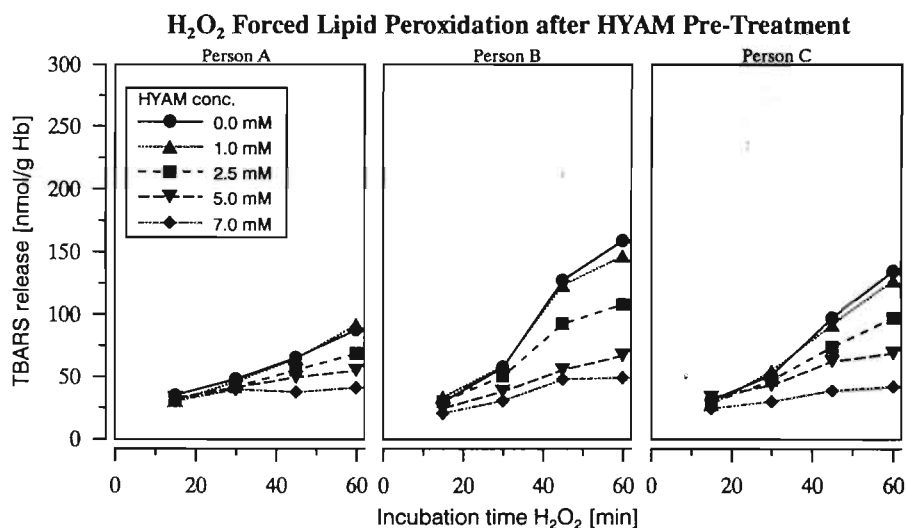


Figure 5.5: Hydrogen peroxide (10 mM) forced lipid peroxidation, expressed as malondialdehyde equivalents in reactivity towards TBA, in erythrocytes of three individuals after pre-incubation of blood with various concentrations of HYAM (1 hour, 37°C).

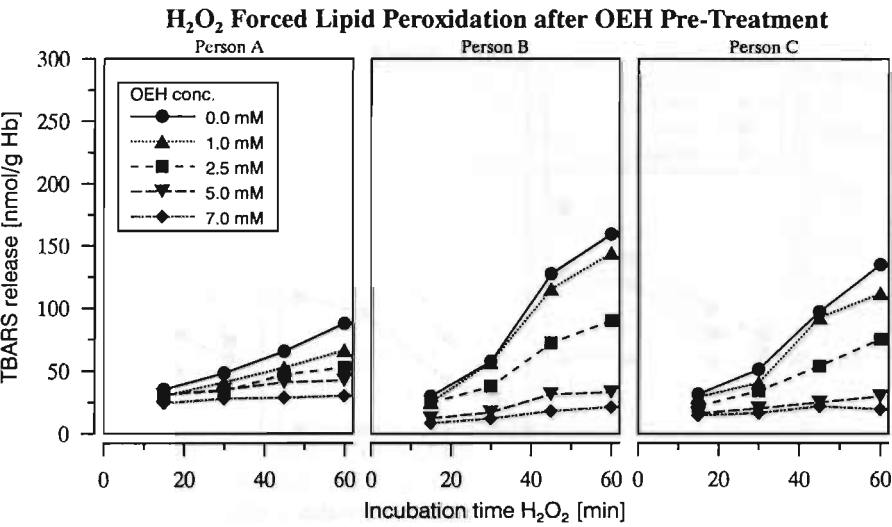


Figure 5.6: Hydrogen peroxide (10 mM) forced lipid peroxidation, expressed as malondialdehyde equivalents in reactivity towards TBA, in erythrocytes of three individuals after pre-incubation of blood with various concentrations of OEH (1 hour, 37°C).

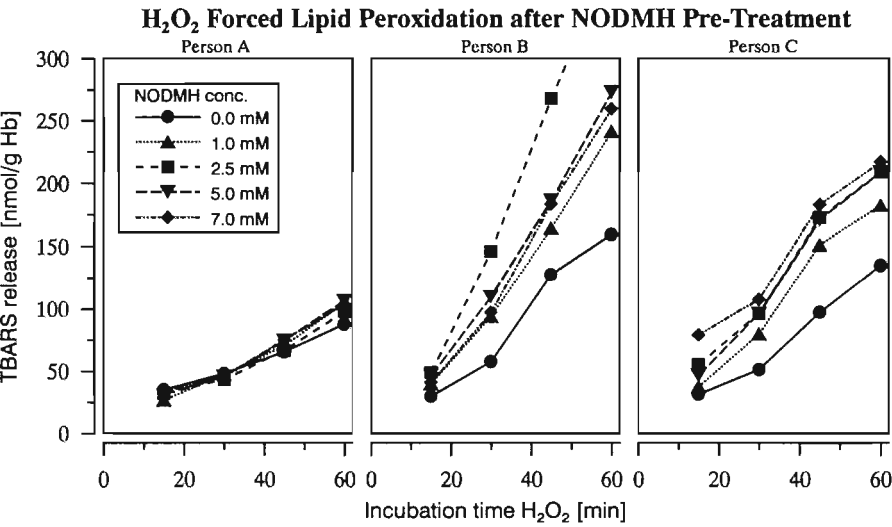


Figure 5.7: Hydrogen peroxide (10 mM) forced lipid peroxidation, expressed as malondialdehyde equivalents in reactivity towards TBA, in erythrocytes of three individuals after pre-incubation of blood with various concentrations of NODMH (1 hour, 37°C).

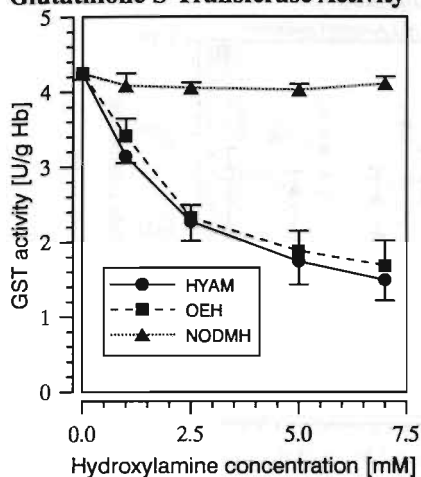
Glutathione S-Transferase Activity

Figure 5.8: Activity of glutathione S-transferase in human erythrocytes after incubation with hydroxylamines (1 hour, 37°C). Regression analyses showed significant decreases in GST activity with increasing HYAM and OEH concentrations ($p < 0.0001$).

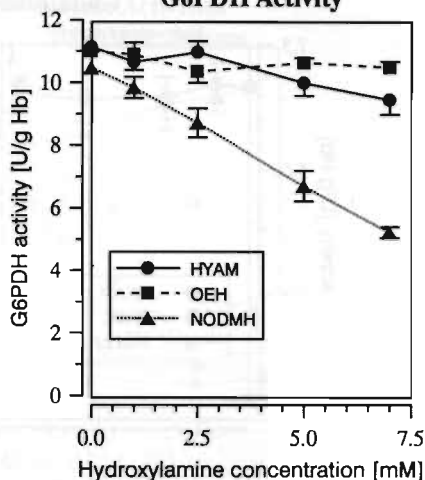
G6PDH Activity

Figure 5.9: Activity of glucose 6-phosphate dehydrogenase in human erythrocytes after incubation with hydroxylamines (1 hour, 37°C). Regression analyses showed significant decreases in activity of G6PDH with increasing concentrations of OEH and NODMH ($p < 0.0001$).

Hb⁻¹lost/mM NODMH), leading to about 50% reduction at the highest (=7 mM) NODMH concentration tested (figure 5.9). For OEH a much smaller decrease was found (0.23 (SE 0.04) U·g Hb⁻¹/mM OEH). For HYAM only a small tendency towards impairment of G6PDH activity was found, which was not statistically significant ($p = 0.14$). The activity of another pentose-phosphate shunt enzyme tested, glucose-phosphate isomerase, was not significantly influenced by any of the three hydroxylamines. NADH-HbR activity was only impaired by HYAM. The control activity of 1.9 U·g Hb⁻¹ fell to about 1.5 U·g Hb⁻¹ in 7 mM HYAM incubations (figure 5.11A on page 91). NADPH-HbR activity was strongly decreased in incubations with both HYAM and OEH. For both substances the decrease amounted to about 0.17 U·g Hb⁻¹ for each mM addition, and a result less than 20% of the control activity was found in the 7 mM incubations (figure 5.11B on page 91).

Discussion

On the whole the effects caused by HYAM and OEH were very much alike, while the effects of NODMH differed markedly. A schematic representation of the mechanisms involved is given in figure 5.12 on page 92. For HYAM and OEH massive

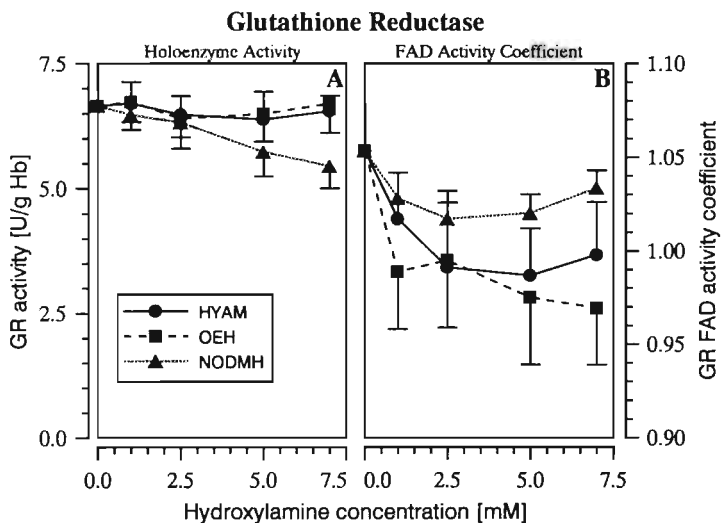


Figure 5.10: A: Activity of glutathione reductase (holoenzyme) and B: FAD activity coefficient of glutathione reductase in human erythrocytes after incubation with hydroxylamines (1 hour, 37°C). The coefficients were calculated by division of the activity in the presence of FAD (11.75 μ M) by the activities determined in a parallel assay without FAD addition. Regression analyses showed a significant decrease in GR activity with increasing NODMH concentration ($p = 0.0002$) as well as significant decreases in FAD activity coefficients with increasing HYAM ($p = 0.025$) and OEH ($p = 0.010$) concentrations.

methemoglobin formation was found, for NODMH this was much lower. HYAM forms complexes with hemoproteins, for instance leading to inactivation of catalase [1, 16, 42]. Methemoglobin formation *in vitro* was previously described for HYAM [7, 11, 16, 18, 63, 66] and NODMH [11]. The mechanism of methemoglobin formation by HYAM is known to differ from that induced by nitrite [18]. Reaction of HYAM with hemoglobin involves Hb/Hb⁺ cycle reactions [7, 16], and leads to decomposition of HYAM. The methemoglobin forming reaction, which produces NH₃ from HYAM, was found to be very fast, explaining the high methemoglobin concentrations found. During the other step of the cycle N₂ is formed. Stolze and Nohl [63] proved with electron spin resonance spectroscopy techniques that free hydronitroxide radicals (NH₂O[•]) are formed as intermediates in the methemoglobin formation induced by HYAM. Stoichiometric considerations lead them to the expectation that H₂O₂ and active oxygen species might also be formed. This radical formation can be one of the causes for LP.

The reduced availability of GSH will also facilitate LP. GSH consumption can be the result of direct oxidation, of reversed GR activity as resulting from the need for extra NADP reduction, and of use in the protective reactions catalysed

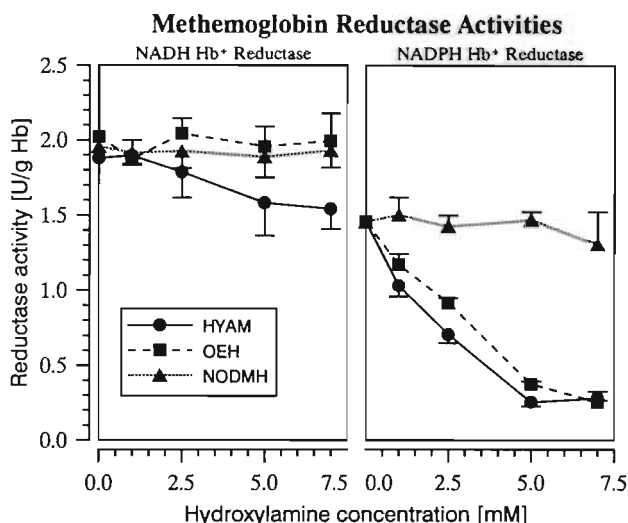


Figure 5.11: Activity of A: NADH methemoglobin reductase and B: NADPH methemoglobin reductase in human erythrocytes after incubation with hydroxylamines (1 hour, 37°C). Regression analyses showed a significant decrease in NADH methemoglobin reductase with increasing HYAM concentrations ($p = 0.002$) and significant decreases in NADPH methemoglobin concentrations with increasing HYAM and OEH concentrations ($p < 0.0001$).

by GST and glutathione peroxidase [15,56]. In this respect LP formation may not only be facilitated by GSH depletion [17] but may also provoke it [56]. This agrees with our finding that OEH as the substance with the highest LP activity also gives the highest GT depletion. Loss of 4-PDS-reactive HbSH groups was not found. After treatment with electrophilic agents such a loss can be clearly demonstrated [23]. However, oxidative loss of HbSH groups for instance by formation of mixed disulfides with GSH, will probably not be picked up by this assay, as dithiodipyridines are able to break GSH sulfhydryl bindings [13]. Such a formation of mixed disulfides may well have occurred in our experiments, since the increases in extracellular GT (mainly present as oxidized glutathione) did not make up for the loss of GT from the erythrocytes. Formation of mixed disulfides from proteins and GSH are a general result of glutathione oxidation [53]. Grossman *et al.* [31] found that mixed disulfide formation in erythrocytes was one of the most important effects of the hydroxylamine metabolite of dapsone. The finding by Kotsifopoulos [43] that HYAM is able to change the electrophoretic behavior of erythrocyte membrane proteins, also indicates the occurrence of mixed disulfides.

Hb⁺ formation, GT depletion and membrane damage—as indicated by LP—are all strongly associated with cellular damages like Heinz body formation

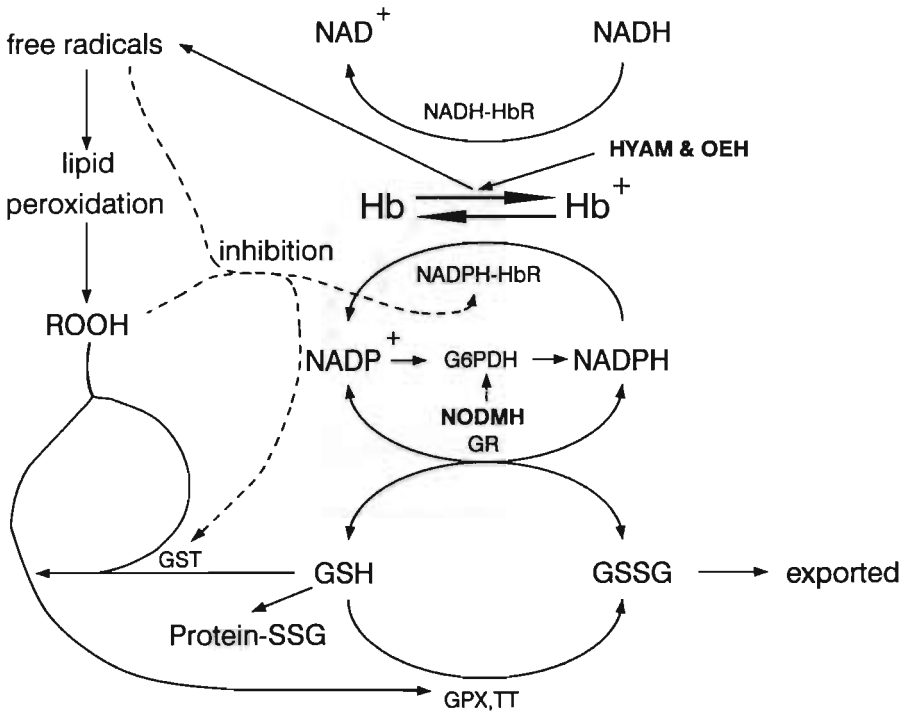


Figure 5.12: Schematic representation of the proposed mechanisms for hydroxylamine erythrotoxicity. Hydroxylamine (HYAM) and the O-ethyl derivative (OEH) induce formation of methemoglobin. During this process free radicals are formed that induce lipid peroxidation. The free radicals plus the lipid peroxides (ROOH) formed lead to inhibition of glutathione S-transferase (GST) and NADPH methemoglobin reductase (NADPH-HbR). The oxidative stress occurring leads to depletion of glutathione (GSH) by various routes. N-substituted hydroxylamines like N,O-dimethyl hydroxylamine (NODMH) inhibit glucose 6-phosphate dehydrogenase (G6PDH), and to a lesser extent glutathione reductase (GR). The resulting inability to reduce NADP^+ and glutathione renders the erythrocyte more vulnerable to oxidative stress, including that caused by HYAM and OEH. Other abbreviations used: GPX = glutathione peroxidase, Protein-SSG = protein-glutathione mixed disulfides, TT = thiol transferase.

and hemolysis. Induction of this kind of damage by the two substances with high potencies for Hb^+ formation, GT depletion and LP is therefore in accordance with expectations. For HYAM Heinz body formation in mice [69] and in pigs [59] were previously reported.

It is well known that LP products can give rise to inhibition of several enzymes [48, 61, 70], and to protein damage in general [21]. Purified rat GST-P

(7-7), which is equivalent to human GST π — the form of GST predominating in human erythrocytes [28] — is very vulnerable to oxidative stress [47, 57, 58, 68]. The human enzyme itself is also inhibited by H_2O_2 [57]. This was confirmed in control experiments where erythrocytes were treated directly with H_2O_2 (data not shown). Previously we showed that GST activity in human erythrocytes is impaired by occupational exposure to coal tar products [24] and to the pesticide dichloropropene [10], in miners with early forms of pneumoconiosis [26] and after long-distance running [25]. At least in the latter two cases oxidative stress is the most likely cause for GST activity loss. Therefore, activity loss of GST under conditions causing LP was expected beforehand. In fact these experiments were included in order to examine the possible use of GST activity loss as a biomarker for hydroxylamine exposure. As expected, GST inactivation was only found for the strong LP causing substances HYAM and OEH and not for NODMH. NADPH-HbR was also strongly inhibited by HYAM and OEH and not by NODMH. Inhibition of NADPH-HbR by HYAM was previously shown by Layne and Smith [44], and does explain the lowered effectiveness of methylene blue treatment in case of HYAM poisoning in mice [60]. The absence of inhibition by NODMH indicates a possible relation between radical stress and inhibition of this reductase. The loss of riboflavin, as indicated by the decreases in GR_{coeff} after HYAM and OEH treatment, might also be caused by consumption during radical scavenging.

NODMH showed far less activity in most aspects discussed so far. Hb^+ formation was low, LP, hemolysis and Heinz body formation were not found, GST and NADPH-HbR inhibition were absent, and riboflavin loss was not found. On the other hand NODMH treatment of erythrocytes resulted in a strong decrease of G6PDH activity, consistent with the earlier findings by Martin *et al.* [46]. GR activity was also decreased by NODMH, but since more than 80% of GR activity was still present at NODMH concentrations of 7 mM, this may be of less importance. Interestingly GT concentrations in erythrocytes were decreased by NODMH while GT in plasma was even more increased than for the two other hydroxylamines. GT depletion by NODMH can be the result of oxidation in glutathione peroxidation and thiol transferase reactions, combined with lower availability of NADPH reduction equivalents for GR activity, due to G6PDH inhibition. Even the G6PDH inactivation alone will render the erythrocyte much more vulnerable to oxidative stress from subsequent exposure to other oxidative compounds. This is confirmed by the fact that NODMH treated erythrocytes did show a higher vulnerability towards subsequent H_2O_2 induced LP. The same kind of effect was to be expected if a glucose-phosphate isomerase inhibition had occurred [55], but such an inhibition was not found for any of the hydroxylamines tested. Contrary to what might have been expected HYAM and OEH pretreatment did not lead to an increased vulnerability to H_2O_2 . This is however easily explained. For H_2O_2 induced LP in erythrocytes ferrous-hemoglobin is a prerequisite [14]. After HYAM and OEH treatment a large fraction of hemoglobin is

converted to the ferric form, necessarily leading to a decrease in LP. It should be noted that changes in H_2O_2 resistance cannot be the result of catalase inhibition by the hydroxylamines, since NaN_3 was added to the assays in order to obtain a full inhibition of catalase. HYAM is known to inhibit catalase both *in vitro* [1,16] and *in vivo* in mouse liver [42], and this inhibition will in itself increase the vulnerability of erythrocytes to oxidative stress.

To our knowledge induction of LP by aliphatic hydroxylamines has not been described previously. The occurrence of this phenomenon in erythrocytes exposed to HYAM and OEH indicates the availability of **free radicals**, possibly including active oxygen species. The presence of active oxygen species and of the resulting peroxides *in vivo* is associated with increased cancer risks and accelerated aging [4,19,27,40,64,67]. It should be noted however that radical production in this case seems to be strongly linked to the presence of hemoglobin. In pilot experiments with HYAM we did not see any LP in washed erythrocyte membranes or in isolated rat hepatocytes. Since erythrocytes do not contain a nucleus, direct DNA damage is excluded. On the other hand reactive products formed in erythrocytes might of course reach other cells and provoke damage there, and sortlike radical mechanisms might be induced by hydroxylamines at hemoproteins elsewhere. In the carcinogenicity studies performed so far [34,49,50,72], no clear *carcinogenic effects* were found. However, the *quality of these studies is insufficient* to decide whether HYAM is an animal carcinogen [8]. HYAM showed some anti-carcinogenic activity with respect to virus induced mammary tumors in mice when treatment was started at an early age [49]. Some other carcinostatic effects of hydroxylamine have also been described [30]. Diethyl hydroxylamine has relatively strong antitumorigenic properties [36,37], it is probably able to terminate free radical processes by sequestration of secondary free radical intermediates [52].

Considering the above, the most serious threat resulting from possible exposure of humans to hydroxylamines seems to come from the direct erythrotoxic effects. Hb^+ formation, LP, GT depletion and inhibition of protective enzymes can lead to faster senescence of erythrocytes and increased sequestration in the spleen. Next to clinical parameters for anemia and increases in reticulocytes, determination of GST and NADPH-HbR and assessment of LP *in vivo* [51] may be useful biomarkers to monitor human exposure to hydroxylamines with direct oxidative activity, like HYAM and OEH. Two special aspects should be noted: 1) due to the inhibition of NADPH-HbR by hydroxylamines like HYAM and OEH and due to the inhibition of G6PDH by compounds like NODMH and trimethyl hydroxylamine, treatment of accidental methemoglobinemia with methylene blue may fail, 2) exposure to compounds like NODMH and trimethyl hydroxylamine will inhibit G6PDH and thereby will render the erythrocytes more vulnerable to a subsequent exposure to direct oxidative compounds. Because of the latter, determination of G6PDH activities may also serve as a valuable biomarker when exposure to this group of hydroxylamines can occur.

Acknowledgment. The authors are grateful to Dr. L. Volovics and Dr. A. Kester of the department for Methodology and Statistics for valuable advices on statistical procedures, to Astrid Feijts for assistance with the analyses of glucose-phosphate isomerase and NADPH-HbR; to Mrs. M. Catsberg from the "Maasland Hospital" (Sittard, NL) for her assistance with the Heinz body determinations and to the "Red Cross Blood-bank, South-Limburg" for the blood samples used. Financial support was given by DSM Special Products b.v. (Geleen, NL).

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Chapter 6

Biological Effect Monitoring

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Published in: *Arch Toxicol* suppl. 15, 1992, 268–277.

Introduction

Monitoring activities have an important role in occupational and environmental practice. Techniques to detect and to control chemical exposure in order to protect people from environmental and occupational illness are rapidly expanding. Most obvious are the methods related to the evaluation of the presence of xenobiotic chemical agents either in the contaminated environment or in the exposed organism. In the first case the procedures are indicated as *environmental monitoring* (EM). In inhalatory exposure, estimations can be made of the 'intake' or 'external load' on the basis of known concentrations in the ambient air. The actual 'uptake' or 'internal load', however, can only be approximated by means of EM methods. On the other hand, measurement of the chemical agents and/or their metabolites in tissues, secreta or excreta of the exposed organism provides an estimate of the total uptake. The latter approach has been defined *biological monitoring* (BM) (Berlin 1984). In addition to these methods many other preventive activities have

Table 6.1: Overview of monitoring methods in occupational and environmental toxicology.

	Monitoring type:	Assessment of:
EM	Environmental Monitoring	external exposure
BM	Biological Monitoring	internal load
BEM	Biological Effect Monitoring	biological effects (not necessarily health related)
HS	Health Surveillance	health state

been developed, including periodic medico physiological examinations of the exposed people. This approach is more directly related to the detection of early health effects, and is defined as *health surveillance* (HS). Changes in physiological functions of the organism can be indicative of changes in the state of health as a consequence of a certain chemical exposure. Such information on the state of health or early impairment of health is very useful in the prevention of overt intoxications due to continuation of overexposures. Well-known examples in this area are the increased levels of serum transaminases providing information on hepatotoxic effects, e.g. after exposure to certain halogenated alkanes. Most of the HS parameters a priori have to be regarded as adverse effects, which means that their significance in terms of prevention is limited. 'Early' adverse effect should also be prevented. Generally, an impaired state of health of the organism can be considered as an endpoint of a sequence of several events at lower levels of homeostasis. Such a 'causality chain' can be depicted as shown in figure 6.1.

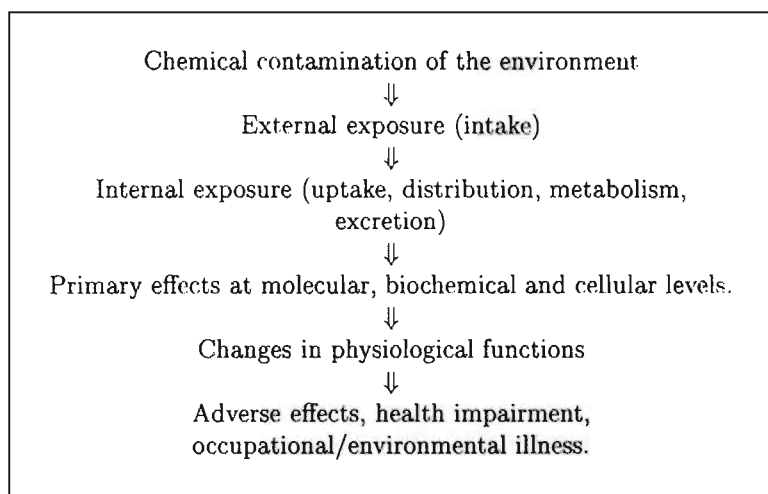


Figure 6.1: Causality chain leading to health effects.

In recent years the arsenal of monitoring techniques has been extended with test methods directed to the assessment of early, possibly reversible, biological effects (*Biological effect monitoring: BEM*). Mostly it is unknown a priori whether primary biological effects have to be regarded as adverse or nonadverse. In this respect the health significance of biological effect parameters has still to be established. In order to avoid confusion among the various monitoring programmes, Zielhuis and Henderson (1986) proposed the following definition:

Biological effect monitoring (BEM) is “the measurement and assessment of early biological effects, of which the relationship to health impairment has not yet been established, in exposed workers to evaluate exposure and/or health risk compared to an appropriate reference”.

Most of the tests for EM and BM are agent specific or specified for certain groups of related chemical, e.g. thioethers, mutagens. The ultimate objective of EM and BM is to estimate the target dose. Because this dose or concentration in the critical organ is determined by toxicokinetic processes, (distribution, biotransformation, elimination), BM is based on ‘the impact of organism on agent’. On the other hand, effect parameters (BEM, HS) refer to toxicodynamic processes and reflect the impact of the target dose on the state of health. Biological effects may be due to many causes: chemical, nonchemical as well as multifactorial causes may be involved. Therefore, BEM and HS are generally agent nonspecific.

Example: Effects on the Glutathione Dependent Detoxification System in Erythrocytes

Some aspects of the BEM approach can be exemplified by methods that are based on changes in the glutathione system in erythrocytes after exposure to electrophilic agents.

Glutathione Depletion

Depletion of liver glutathione (GSH) during exposure to chemically reactive compounds is a well-known phenomenon both *in vitro* and *in vivo*. GSH is also involved in the metabolism of many substances that pose an occupational health hazard. This leads to the urinary excretion of thioethers (van Doorn *et al.*, 1981; Henderson *et al.*, 1984), which itself is used for biological monitoring. In principle, exposure to such compounds might lead to a decrease in GSH concentrations not only in the liver but also in more easily accessible material like blood cells. *In vitro* GSH in erythrocytes can be depleted with thiol reactive substances like iodoacetamide, N-ethylmaleimide and diethylmaleate (Evelo and Henderson, 1988) and also by metabolites of 3-acetamidophenol and cyclophosphamide (Palmen and Evelo, 1993). The reactivity of GSH is so high that it results in an effective protection of cellular macromolecules. Figure 6.2 shows that during *in vitro* treatment of erythrocytes with iodoacetamide and N-ethylmaleimide the hemoglobin cysteine is not significantly alkylated as long as the intracellular GSH is not fully depleted. This kind of observations prompted us to study GSH levels in some situations where people are potentially exposed to electrophilic agents.

Glutathione S-Transferase Inactivation

The activity of erythrocyte glutathione S-transferase (GST ρ) is another interesting, and biochemically related, parameter. The activity of GST in general is known to be susceptible to many chemicals (Mannervik and Danielson, 1988). The *in vitro* susceptibility of GST activity from human erythrocytes to industrially important electrophiles like acrolein, propylene-oxide, styrene oxide, ethylene dibromide and ethylene dichloride has been described by Ansari *et al.* (1987). Kilpikari and Savolainen (1984) reported decreased values of erythrocyte GST activity in workers exposed to hot rubber fumes. Normal GST activity is known to vary largely between individuals (Strange *et al.*, 1982). In this respect the comparison of individual pre- and post exposure values is expected to be more sensitive than the comparison with a control group.

Creosote Workers

In a pilot study we determined GST activities in erythrocytes and GSH in blood of two workers creosoting wood. The samples for this study were collected as part of a biomonitoring study carried out by Jongeneelen and co-workers. Blood and

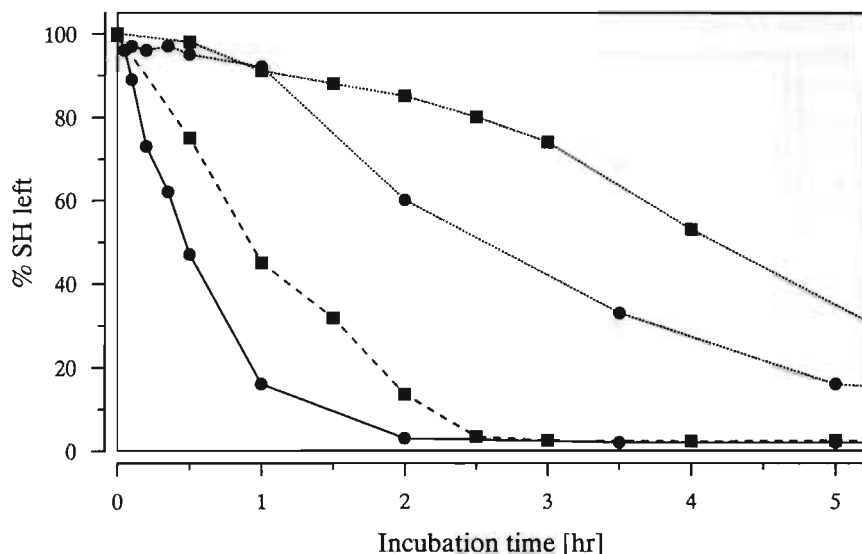


Figure 6.2: Free sulfhydryl groups of hemoglobin (Hb,) and glutathione (GSH, —) after incubation of human blood with the depletors iodoacetamide (IACa, ●, 1 hour) and N-ethylmaleimide (NEM, ■, 15 min.) at 37°C. Data are the mean of duplicate determinations. 100% HbSH values were 2.15 and 2.18 moles/mole and 100% GSH values were 0.86 and 0.72 mM respectively. Determined by titration with 4,4'-dithiodipyridine (HbSH) and 5,5'-dithiobis 2-nitrobenzoic acid (GSH).

urine samples of two exposed workers and four control persons, all nonsmokers, were collected: at the beginning of a work week—which was after an exposure free period of several weeks—at the end of that week and after the weekend. Total (i.e. reduced plus oxidized) glutathione was determined with the cyclic oxidation reduction method (Anderson, 1985). GST activity towards 1-chloro-2,4-dinitrobenzene (CDNB) was determined essentially as described by Habig and Jakoby (1981). Both methods were slightly modified as described previously (Brouwer *et al.*, 1991). Figure 6.3 shows the separate GSH and GST values for the two workers. In order to exclude errors caused by day to day analytical variations, values were expressed as the ratio of the individual value and the mean value of the four control persons determined the same day. In both workers GST and GSH values were decreased at the end of the week, and returned to values close to the initial values after the weekend. These data, which in themselves have a rather limited value, become more interesting when related to some other parameters determined in the same workers, as described below.

Environmental monitoring of pyrene and biological monitoring of urinary 1-hydroxypyrene, as markers for the exposure to polycyclic aromatic hydrocarbons,

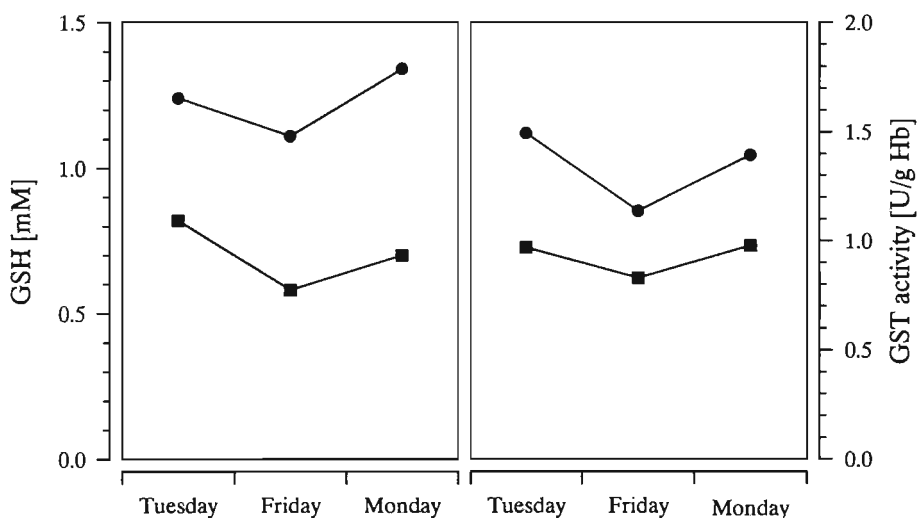


Figure 6.3: Glutathione concentrations in blood and Glutathione S-Transferase activities in erythrocytes of two workers (● and ■) creosoting wood. Samples were collected before and after the work period and after the weekend. All values are the result of determinations in duplicate, and were expressed as relative values compared to samples from four nonexposed persons collected and determined on the same day.

were done by Jongeneelen and coworkers (Dr. F. Jongeneelen, personal communication). Environmental monitoring revealed that dermal and inhaled exposition for person 1 were generally at least two times higher than for person 2. Hydroxypyrene in urine was determined twice each day with previously described methods (Jongeneelen *et al.*, 1987). The individual excretion values for hydroxypyrene on the morning of the sampling days for the blood samples are shown in table 6.2. For both workers the urinary hydroxypyrene values were clearly above the normal range (normal 95 percentile value is $0.66 \mu\text{mol/mol}$ creatinine) at the end of the week. The interindividual differences in excretion were in accordance with the BM data.

Even more interesting were the findings by Steenwinkel and Baan (Dr. R. Baan personal communication) who determined DNA adducts of polycyclic aromatic hydrocarbons in white blood cells with the ^{32}P -postlabeling method developed by Randerath and co-workers (Gupta *et al.*, 1982, Randerath *et al.*, 1981; Watson, 1987) in the same blood samples used for the GSH and GST determinations. The results are listed in table 6.2. All values are the means of two, in some cases three, determinations. The analytical variation between experiments was about 20%. The end of week values are clearly elevated compared to the

pre-exposure values. For the DNA adducts a tendency for return to normal values is visible too, but this effect is small compared to the analytical variability. A slowness in recovery for DNA alkylation need not only be the result of slow repair processes, but can also be caused by continued availability of reactive compounds from internal storage in for instance fat tissue. Faster recovery of GSH and GST would, on the other hand, not only indicate the relatively fast reversibility of their impairment, but would also indicate that this impairment is caused by substances whose residency is relatively short.

Table 6.2: DNA adduct values in creosote workers.

Sampling day	DNA adducts [attomoles/ μ g DNA]		Hydroxypyrene exc. [μ moles/mol creat.]	
	person 1	person 2	person 1	person 2
Tuesday	6	n.d. ^a	0.35	0.70
Friday	15	19	2.8	1.15
Monday	13	18	0.55	0.5

a. n.d. is not detectable, detection limit was 4 attomoles/ μ g DNA.

Dichloropropene Workers

Further indications for the usefulness of the determination of GSH and GST came from a prospective study which was performed in the Dutch flower bulb culture, in collaboration with Brouwer and de Wolff, to investigate the possible effects of the soil fumigant 1,3-dichloropropene (DCP) (Brouwer *et al.*, 1991). At the start of the season in July, and after the season in October, blood and urine samples of workers applying DCP were collected. The activity of erythrocyte GST was significantly ($p < 0.002$; Wilcoxon matched pair analysis) decreased from 4.7 before to 3.3 U/g Hb after the season. The GSH values in blood were also significantly ($p < 0.02$) decreased from 0.93 to 0.82 mM. With a single exception for an unusual low GSH value in the pre-season samples, all individual values were found to be decreased at the end of the season (Fig. 6.4).

The study was combined with environmental and biological monitoring of 1,3-dichloropropene (van Welie *et al.*, 1991). In the biological monitoring study excretion of two mercapturic acid metabolites of DCP was determined. Since these mercapturic acids are the result of GSH conjugation, their excretion indicates that GSH is actually used in the metabolism of DCP. Changes in health effect parameters for liver and kidney function in these workers were determined by Verplanke. Decrease of total serum bilirubin together with an increase in serum γ -glutamyltranspeptidase activity indicated moderate hepatic enzyme induction. And a possible subclinical nephrotoxic effect was indicated by significant increases

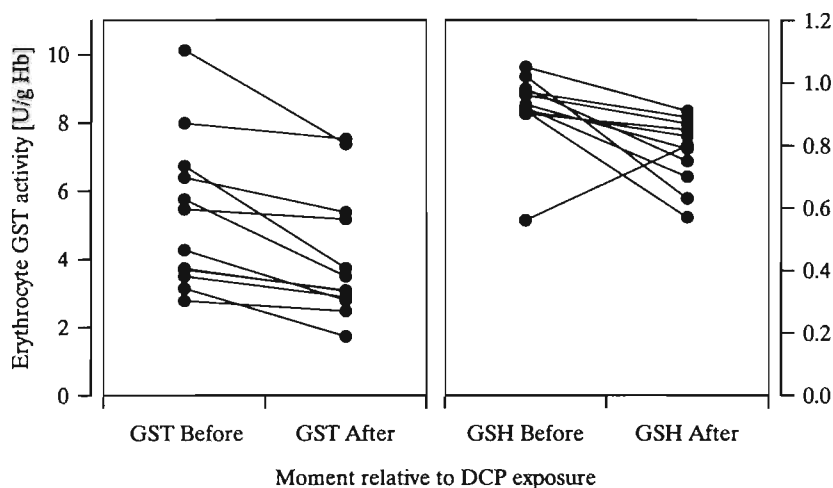


Figure 6.4: Glutathione S-transferase activity in erythrocytes and glutathione in blood of workers exposed to 1,3-dichloropropene.

in urinary excretion of albumin and retinol binding protein. The direction of the change in the determined health effect parameters for the liver —enzyme induction, not liver impairment— and the size of the changes for the kidney functions —within the normal clinical range— lead to the conclusion that there were no significant health effects on the individual level. Nevertheless, significant biological effects were found, which lead to the identification of internal exposure, and possibly increased risk.

Smokers

Since smokers are continuously exposed to electrophilic compounds originating from cigarette smoke, some effects on the glutathione system could be anticipated. Table 6.3 shows the concentration of total GSH in blood, erythrocyte GST activity and the amount of GST present in the erythrocytes. The latter amount was determined by a competitive ELISA assay with antibodies towards human GST π class enzymes (Medlabs, Dublin Ireland).

The thioether excretion in smokers was significantly increased. Such an increase is usually found in smokers (Henderson *et al.*, 1984), and it is included in the table only because it indicates exposure. Total blood GSH did not differ between smokers and nonsmokers. The decrease in erythrocyte GST activity towards CDNB was not statistically significant. The amount of GST π class protein however, was strongly increased in the smokers. This means that the specific activity of the GST π protein present in the erythrocytes of smokers towards the common substrate CDNB is lower than normal. This may indicate that GST

Table 6.3: GST and GSH in blood and urinary thioether excretion of smokers and nonsmokers. Median values and ranges are shown. Symbols indicate significant differences from the values for nonsmokers by Wilcoxon 2-sample test; *: $p < 0.10$; ‡: $p < 0.02$.

	nonsmokers	smokers
age (mean ± s.d.)	33 ± 7 n=11	37 ± 10 n=10
GSH [mM]	1.06 0.88–1.32	1.12 0.77–1.17
GSTρ act. (CDNB) [U/ml blood]	0.71 0.42–1.13	0.52 0.31–0.95
GSTπ conc. (ELISA) ≡[U/ml blood]	0.6 0.3–4.5	4.2‡ 1.6–5.0
Thioether excr. [mmol/mol creat.]	3.3 0.5–9.9	4.7* 3.0–14.0

synthesis during erythrocyte formation is induced in smokers, to compensate for inactivation that takes place during the erythrocyte life cycle. The induction of a GST not active towards CDNB would be another explanation.

Physical Activity

Since part of the GST activity loss may be caused by reactive oxygen species, we decided to study the influence high oxygen consumption due to physical activity (Evelo *et al.*, 1992). Previously sedentary men ($n = 23$) and women ($n = 18$) were trained to run a 1/2 marathon contest after 40 weeks. After 20 weeks of training total blood GSH and erythrocyte glutathione reductase (GR) activity were increased. After 40 weeks total blood GSH values were returned to normal, while GR remained elevated. Erythrocyte GST activity was strongly increased after 40 weeks, this may be indicative of the occurrence of lipid peroxidation in this training phase.

The participants ran a 15-km race after the first 20 weeks of training and a 1/2 marathon after 40 weeks. After the 15-km race total blood GSH and erythrocyte GST activity were decreased, while GR activity was increased. GSH and GR values were returned to before race conditions 5 days after the 15-km race. After the 1/2 marathon total blood GSH and erythrocyte GR activity were increased, while GST activity was decreased. 5 Days after the 1/2 marathon all values were

returned to before race conditions. The difference in GSH sensibility after both races may be caused by differences in before race conditions. GSH was elevated before the 15-km race. The increases in GR activities after the races could at least partially be explained by concurrent increases in riboflavin concentrations.

Concluding Remarks

More and more methods in occupational and environmental toxicology, allow the determination of biological effects of which either no clear relationship to health impairment has been established, or at levels where no known clinical significance exists. The primary value of the use of such methods lays in the relation between the determined effects and the internal dose. If the relation between internal dose and the ultimate risk is known, like it is suggested for hemoglobin and DNA adducts (Henderson *et al.*, 1989), the assessment of biological effects can be used for risk estimations. However, even if such a clear relationship is not established the determination of biological effects may be indicative of exposure to reactive compounds, and may be used for the control of exposure situations. Much opposition against the use and development of BEM methods is based on a bad understanding of their results. Problems arise when health significance is attributed to results which do not allow such conclusions. It has often been overseen that methods leading to such results can still be very valuable for the management of risk. A clear distinction between BEM and HS is needed to improve this situation.

The development of new BEM methods depends largely on knowledge of toxicodynamics. The development of the methods for the glutathione system for instance, was stimulated by the understanding of the biochemical processes leading to thioether excretion and by insight in reactive oxygen toxicity. To put it in another way, knowledge of toxicodynamics can, and in our view should, be used for the development of new methods for biological effect monitoring.

Acknowledgements. The authors wish to thank Dr. Jongeneelen from the university of Nijmegen and Dr. Baan from the Medical Biological Laboratory of TNO for the opportunity to reference unpublished data on hydroxypyrene excretion and DNA adducts in the creosote workers.

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Chapter 7

Biological Effect Monitoring of Occupational Exposure to 1,3-Dichloropropene: Effects on Liver and Renal Function and on Glutathione Conjugation

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Published in: *Br J Ind Med* 48, 1991, 167-172*.

*. A reaction to the original paper by N.J. van Sittert, G.E. Veenstra, E.P. Dumas and W.F. Tordoir from Shell International and a reply by the authors was published in: *Br J Ind Med* 48, 1991, 646-648.

Summary. A prospective study was performed in the Dutch flower bulb culture to investigate the possible effects of subchronic exposure to the soil fumigant 1,3-dichloropropene (*DCP*) on liver and kidney function and on glutathione conjugation capacity in blood. Urine spot samples and venous blood samples from 14 workers applying DCP (applicators) were taken at the start of the season in July, and after the season in October. The parameters of liver function measured were: alanine aminotransferase, aspartate aminotransferase, alkaline phosphatase, lactate dehydrogenase, gamma-glutamyl transpeptidase, and total bilirubin (conjugated and unconjugated). Total bilirubin was significantly decreased from 9.5 before to 7.0 μM after the season. In combination with an increase in serum gamma-glutamyl transpeptidase activity from 12.5 to 19.5 $\text{U}\cdot\text{l}^{-1}$, this indicates moderate hepatic enzyme induction. To study renal function, creatinine and β_2 -microglobulin in serum, and β_2 -microglobulin, albumin, alanine aminopeptidase, γ -galactosidase, and retinol binding protein in urine were measured. The glomerular function parameters albumin in urine and creatinine in serum changed significantly during the season: albumin concentration increased from 5.2 to 7.6 $\text{mg}\cdot\text{l}^{-1}$, whereas creatinine excretion decreased from 93.0 to 87.5 μM . The tubular function parameter retinol binding protein also increased in concentration from 20.0 to 26.9 $\mu\text{g}\cdot\text{l}^{-1}$. Therefore, a subclinical nephrotoxic effect of subchronic exposure to DCP cannot be excluded. Effects on glutathione conjugation capacity were studied by measuring erythrocyte glutathione S-transferase activity and blood glutathione concentrations. The activity of glutathione S-transferase in erythrocytes was significantly decreased from 4.7 before to 3.3 $\text{U}\cdot\text{g Hb}^{-1}$ after the season. The same was true for the blood glutathione concentrations, which decreased from 0.93 to 0.82 mM. Both parameters seem to be affected by exposure to DCP.

Introduction

A mixture of Z- and E-1,3-dichloropropene (*DCP*) is used extensively in the Dutch flower bulb culture for soil fumigation. This is mainly performed by commercial applicators. Products containing DCP, such as Telone II and Shell-DD 95, are injected into the soil, without previous dilution, in the period between June and November. During field application of soil fumigants containing DCP, applicators may be exposed to DCP, mainly by inhalation.

The primary target organs of DCP toxicity in experimental animals are the liver and the kidney. Exposure of rats and guinea pigs to 50 ppm (220 $\text{mg}\cdot\text{m}^{-3}$) DCP vapor by inhalation for seven hours a day, five days a week during one month induced liver and kidney injury. Exposure to 3 ppm (13.6 $\text{mg}\cdot\text{m}^{-3}$) for six months induced cloudy swelling of the renal tubular epithelium in male rats. This was found to be reversible after stopping exposure [22]. Based on evidence of no effects in rats, mice, guinea pigs, and dogs exposed seven hours a day to 1 ppm

($4.5 \text{ mg} \cdot \text{m}^{-3}$) for 6 months, and of the reversible injury in male rats exposed to 3 ppm, a time weighted averaged threshold limit value (*TLV*) of 1 ppm has been recommended for dichloropropene [1]. This is also the present Dutch occupational exposure limit (*OEL*) [16].

Apart from a recent study about short-term exposure to DCP and renal effects [18], no studies of exposure to DCP on health effects in man have been published.

In combination with an environmental and biological monitoring study of DCP in the Dutch flower bulb culture [6,26], the present prospective exploratory study was carried out to investigate the early effects of subchronic exposure to DCP on liver and renal function of commercial applicators.

Exposure to electrophilic compounds such as DCP or its metabolites might lead to a loss of biologically active thiol groups and assessment of this loss could be useful as a biological marker of exposure [10]. In this study two parameters of change in thiol groups — namely, erythrocyte glutathione S-transferase activity and blood glutathione concentration — were also evaluated.

Methods

Population and study design

The study was restricted to the "Bollenstreek", the area between Leiden and Haarlem (The Netherlands) where the flower bulb culture is mainly concentrated. All 14 commercial applicators using DCP in soil fumigation in this region participated in the study, which was approved by the medical ethics committee of the University Hospital of Leiden.

By means of a questionnaire, completed in July on the first sampling day, information was obtained from each subject on age, smoking and drinking habits, medication, and current and past diseases. The mean age was 42 years (median 40, range 33–60). Five workers smoked, with an average rate of 15–20 cigarettes a day (range 1–5 to >20). The average alcohol intake was 5–10 glasses a week (range <1 to >60). One applicator did not drink alcoholic beverages. Two applicators used an oral anticoagulant (phenprocoumon). None of the workers was known to have liver or kidney disease.

For 12 applicators blood and urine samples were collected between 12.00 a.m. and 2.00 p.m. both before and after the season; for two, samples were collected between 6.00 and 7.00 p.m. before and after the season.

Possible effects of exposure to DCP on the liver were studied by measuring alanine aminotransferase (*ALAT*), aspartate aminotransferase (*ASAT*), lactate dehydrogenase (*SLDH*), γ -glutamyl transpeptidase (γ *GT*), and alkaline phosphatase (*ALP*) activities, and total bilirubin concentrations (*Bil-tot*) in serum. To study effects on renal function, parameters were chosen to discriminate between effects on glomerular and tubular function. Glomerular function was measured

by assessing concentrations of β_2 -microglobulin in serum (β_2 -M-S), creatinine in serum (*Creat-S*), and albumin in urine (*Alb-U*). Proximal tubular function was reflected by the excretion of β_2 -microglobulin (β_2 -M-U), retinol binding protein (*RBP*), β -galactosidase (β -gal), and alanine aminopeptidase (*AAP*) in urine. Effects on glutathione conjugation capacity in blood were evaluated by measuring blood glutathione (*GSH*) concentrations and the activity of erythrocyte glutathione S-transferase (*GST*: EC 2.5.1.18).

Analytical methods

Blood samples for measurement of liver and renal function parameters were collected by vein puncture using 10 ml Vacutainer tubes. Blood was allowed to clot and transported within two hours in a cool box to the laboratories for Clinical Chemistry and Toxicology, University Hospital of Leiden. For the determination on GST activities and GSH concentrations, blood samples were collected in 10 ml Vacutainer tubes containing ethylenediaminetetra-acetic acid (*EDTA*). The samples for the determination of ALAT, ASAT, SLDH, γ GT, and ALP activities and Bil-tot and Creat-S concentrations in serum were analyzed within 24 hours with a SMAC multichannel analyzer (Technicon Instruments Corporation, Tarrytown, NY, USA), according to the manufacturer's instructions. Concentrations of the protein β_2 -M in urine and serum were measured with a commercial radioimmunoassay kit (Pharmacia Diagnostics AB, Uppsala, Sweden). For the determination of Alb-U and RBP concentrations and β -Gal and AAP activities, the urine samples were collected and transported on ice to the Coronel Laboratory, University of Amsterdam, prepared the same day, and analyzed within one week. Concentrations of Alb-U and RBP were analyzed with a latex immunoassay according to Bernard and Lauwerys [5]. For the determination of β -Gal activity, a colorimetric assay according to Maruhn [15] was used. Alanine aminopeptidase activity was assayed by the method of Jung and Scholz [11]. In earlier studies reported from the Coronel Laboratory coefficients of variation were: Alb-U 3%, RBP 10%, and β -Gal 5% [23]. Assuming that the ratio between the renal function parameters and the urinary density was the same before and after the season, the data were adjusted for urinary density [23].

Blood samples used for GST and GSH determinations were transported in a cool box to the Department of Toxicology, University of Nijmegen, and prepared (GST) or analyzed (GSH) the same day. Concentration of GSH was determined essentially as described by Anderson [3]. For the determination of GST activity in erythrocytes, red cells were washed three times with phosphate buffered saline (*PBS*; pH 7.0) and lysed by addition of three volumes of water containing 1.4 mM neutral dithiothreitol (*DTT*) to prevent oxidation of thiol groups. After the ionic strength had been restored by adding one volume twofold concentrated PBS containing 1.4 mM DTT, cell fragments were removed by centrifugation at 15,000 *g* for 10 minutes. Only the clear upper part of the supernatant was used for the determinations. The activity towards 1-chloro-2,4-dinitrobenzene (*CDNB*) was

determined as described by Habig and Jakoby [9] with a Cary 119 spectrophotometer in auto-slit mode with the gain adjusted to 2.5.

The concentration of hemoglobin (*Hb*) in the hemolysates was determined with the hemoglobin cyanide procedure [12].

Exposure

After the season, all applicators were asked to estimate the number of hours and days worked in soil fumigation.

Statistical Analyses

To test for significance of differences between values for liver and renal function parameters and blood GST and GSH values measured before and after the season, matched pair analysis was performed with the Wilcoxon non-parametric signed rank test. For β -Gal at pH > 7.0 (5 samples) and for β_2 -M-U at pH < 5.5 (8 samples), data were excluded from the analysis because of instability above or below these pH values [23]. Data were analyzed both with and without adjustment for urinary density.

The Friedman test was used to test for equality between values before and after the season for the liver parameters, and the glomerular and tubular function parameters simultaneously. All statistical analyses were performed with the statistical package of the social sciences SPSS(-X) [17].

Results

Exposure

All 14 applicators took part in soil fumigation with DCP during the season. The total number of working hours for each subject during this period varied from 26 to 300 with a median of 142 hours, divided over 4 to 37 days with a median of 20 days.

The number of hours during which the applicators fumigate soil only indicates the duration and not the level of exposure. These exposure data are not used, therefore, to calculate a dose response relation.

In the environmental monitoring study [6], exposure to DCP for each applicator was measured on one or two days for biological monitoring [26]. These data cannot be considered representative of the average exposure of the different applicators during the season and are, therefore, not suitable for an estimation of subchronic exposure to DCP of the 14 workers in this study.

From the environmental monitoring study it appeared that exposure was comparatively high. The TWA concentration of DCP ranged from 1.9 to 18.9 mg·m⁻³. On 30% of the observed working days the Dutch occupational exposure limit of 5 mg·m⁻³ (1 ppm) was exceeded.

Liver function

With the exception of Bil-tot, no significant differences were found for the parameters of liver function between samples obtained before and after the seasons (table 7.1). Concentration of Bil-tot was significantly decreased after the season from 9.5 to 7.0 μM (median values; $p = 0.025$). Activity of γGT was appreciably increased from 12.5 before to 19.5 $\text{U}\cdot\text{l}^{-1}$ after the season but the difference was only of borderline significance (median values; $p = 0.099$).

Table 7.1: Liver function parameters before and after the season: values corrected for specific density ($n=14$)

	Before season (median (range))	After season (median (range))	p Value
Bil-tot ($\mu\text{mol/l}$)	9.5 (6.0–15.0)	7.0 (1.0–14.0)	0.025*
ALP (U/l)	38.5 (26.0–69.0)	39.5 (26.0–68.0)	0.576
ASAT (U/l)	8.5 (6.0–11.0)	8.0 (5.0–18.0)	0.155
ALAT (U/l)	10.5 (5.0–20.0)	12.0 (6.0–36.0)	0.235
SLDH (U/l)	145.5 (117.0–189.0)	149.0 (119.0–194.0)	0.972
γGT (U/l)	12.5 (5.0–31.0)	19.5 (9.0–34.0)	0.099

*. Statistically significant ($p < 0.05$)

Renal function

For 2 out of 3 glomerular function parameters, Creat-S and Alb-U, significant differences between the concentrations before and after the season were found. Concentration of Creat-S decreased from 93.0 to 87.5 μM (median values; $p = 0.013$)(table 7.2). In one subject, however, Creat-S was very low, especially at the end of the season. This person had been treated with ampicillin one month before the study. Concentration of Alb-U increased from 5.2 to 7.6 $\text{mg}\cdot\text{l}^{-1}$ (median values; $p = 0.044$)(table 7.2). The excretion of albumin in urine is normally less than 8 $\text{mg}\cdot\text{l}^{-1}$ [20]. One applicator had values 3 times higher than this value, before as well as after the season. This person was advised to consult his general practitioner.

The only tubular function parameter that was significantly increased after the season was RBP, from 20.0 to 26.9 $\mu\text{g}\cdot\text{l}^{-1}$ (median values; $p = 0.036$). No significant differences were found for $\beta_2\text{-M}$ concentrations in urine or in serum, and for AAP and $\beta\text{-Gal}$ activities in urine (table 7.2).

The concentration of $\beta_2\text{-M}$ in urine is normally 200 $\mu\text{g}\cdot\text{l}^{-1}$ [20]. Three workers had values above this value but below 300 $\mu\text{g}\cdot\text{l}^{-1}$, which is the normal upper limit according to the manufacturer of the assay kit. A significant difference was

Table 7.2: Renal function parameters before and after the season. Values corrected for specific density (n=14 unless stated otherwise)

	Before season (median (range))	After season (median (range))	p Value
Creat-S ($\mu\text{mol/l}$)	93.0 (74.0–109.0)	87.0 (42.0–100.0)	0.045*
β_2 -M-S (mg/l)	1.3 (0.8–1.8)	1.4 (1.1–1.8)	0.563
β_2 -M-U ($\mu\text{g/l}$)	76.4 (45.0–174.3) (n = 8)	109.9 (65.3–350.9) (n=12)	0.208
AAP (U/l)	9.7 (6.5–13.9) (n = 12)	8.3 (5.0–18.0) (n=13)	0.754
β -Gal (U/l)	1.9 (1.1–7.2) (n = 11)	2.4 (1.7–5.6) (n=12)	0.308
Alb-U (mg/l)	5.2 (1.6–31.8)	7.6 (2.7–37.0)	0.013*
RBP ($\mu\text{g/l}$)	20.0 (8.4–36.1)	26.9 (16.5–72.7)	0.036*

*. Statistically significant ($p < 0.05$)

Table 7.3: Glutathione S-transferase activity (CDNB) in erythrocytes and glutathione concentrations in blood before and after the season (n=12)

	Before season (median (quartile 1–3))	After season (median (quartile 1–3))	p Value
GST (U/g Hb)	4.7 (3.6–6.6)	3.3 (2.8–5.3)	0.002*
GSH (mM)	0.93 (0.90–0.97)	0.82 (0.73–0.86)	0.02*

*. Statistically significant ($p < 0.05$)

found by testing the differences between values before and after the season for all three glomerular function parameters together (median values; $p = 0.011$; Friedman test). Statistical analyses of the data not adjusted for urinary density did not show significant differences in any renal function parameters under study.

Blood GST and GSH

The concentration of GSH in blood decreased significantly from 0.93 before to 0.82 mM after the season (table 7.3; median values; $p = 0.023$). The median GST activity decreased from 4.7 to 3.3 U·g Hb⁻¹ (median values; $p = 0.002$). The GSH concentrations were lowered after the season in all but one person, whereas GST activity was decreased in all persons (figures 7.1 and 7.2).

Lymphocytes from the same blood samples were tested for the presence of

GST [24] The GST activity in erythrocytes and GSH concentrations in blood were not significantly different for the μ positive and μ negative groups.

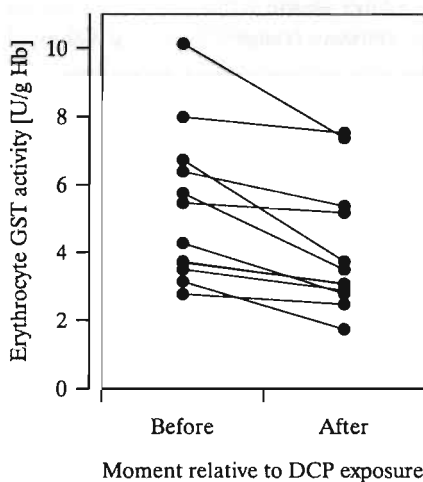


Figure 7.1: Glutathione S-transferase activity in erythrocytes measured before and after the season.

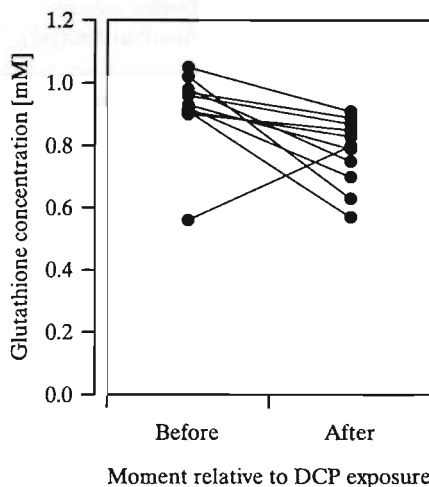


Figure 7.2: Glutathione concentrations in blood measured before and after the season.

Discussion

Monitoring in occupational toxicology is directed towards the assessment of exposure, which may lead to minimization of hazards or towards determination of early health effects, which may serve the added purpose of better risk evaluations [10,27]. The liver and kidney function tests used in this study give information on the functional state of the organs concerned. Changes in GSH and GST may indicate primary biological effects. These cannot be interpreted, as yet, in terms of health effects, but only in terms of endogenous exposure to electrophilic agents. The renal function tests used in this study are considered to be sensitive enough to detect early effects of nephrotoxins and are feasible as indicators of early changes in renal function [14]. Low molecular weight proteins, such as β_2 -M-U and RBP, are freely filtered from the plasma by the renal glomerulus and reabsorbed by the proximal tubular cells. The reabsorption of these proteins by the kidney is almost complete, which explains why their presence in urine is a sensitive indicator of impaired proximal tubular function [14]. In this study a significant increase of Alb-U and RBP in urine was found after the season. If the increase in Alb-U excretion can be attributed to reduction in tubular reabsorption, a much larger increase in RBP excretion would be expected [14]. Both

for Alb-U and RBP the excretion in urine was, however, low compared to other studies.

No significant increase in β_2 -M-U was found, by contrast with what might be expected from another study in which a high correlation between RBP and β_2 -M-U was shown [5]. This may be explained by the small amount of data on β_2 -M-U that could be used in our calculations. Data from 8 out of 28 urine samples had to be excluded from the analysis because of a pH less than 5.5; β_2 -M-U in acid urine is very unstable and the protein degrades quickly.

Increased excretion of high molecular weight proteins such as albumin and ferritin may also result from a reduced tubular reabsorption of the filtered load [14]. An increased excretion of both a high (Alb-U) and a low (RBP) molecular weight protein may indicate a reduced tubular reabsorption capacity in these workers. Effects on the proximal tubules can also be detected by measuring the activity of several enzymes in the urine [14]. Neither of the two urinary enzymes measured in this study, AAP and β -Gal, had significantly increased activities after exposure to DCP during the season. After the present study was completed Osterloh *et al.* [18] described a subclinical effect on renal function expressed in an increased urinary excretion of N-acetyl-glucosamidase after short-term exposure to DCP through soil fumigation. The concentrations exposed to ranged from 0.26 to 9.36 $\text{mg}\cdot\text{m}^{-3}$.

The exposure concentrations measured in our corresponding environmental monitoring study ranged from 1.9 to 18.9 $\text{mg}\cdot\text{m}^{-3}$, and are comparable with those used in the animal studies in which a cloudy swelling of the renal tubular epithelium of male rats was found at 3 ppm (13.6 $\text{mg}\cdot\text{m}^{-3}$) after 6 months of exposure [22]. With regard to liver function, a significant decrease in concentration of Bil-tot in serum within the normal range was found after the season. Together with the remarkable increase in γ GT this may indicate that DCP has enzyme-inducing properties. Induction of the β -glucuronidase activity may result in an increased contribution of conjugated Bil-tot, and hence in an increased elimination. In animal studies, an effect on the liver was also found at a comparable exposure level; in female rats exposed for 6 months to 3 ppm (13.6 $\text{mg}\cdot\text{m}^{-3}$), a higher liver to body weight ratio was found. No histopathological changes in the liver were seen, however, and the increase was considered not to be an adverse effect, although it may be related to exposure [22]. Because the data from the present study were not compared to a control group, the results might be influenced by seasonal variation. When the results were compared, however, with those of a study by Gidlow *et al.* [8] about seasonal fluctuations of some biochemical parameters monitored in industrial workers in the United Kingdom exposed to a variety of potentially toxic chemicals, it appeared that our results (a decrease in creat-S and an increase in γ GT) were opposite to those expected from seasonal fluctuations.

Because of the electrophilic nature of DCP attack on sulfhydryl groups can be expected. Glutathione provides the largest pool of biologically available thiol

groups. Normal GSH concentrations in blood do not differ much between people of the same age group [2].

The *in vitro* susceptibility of erythrocyte GST activity to industrially important electrophilic agents has been described [4]. The authors expected to find induction of GST synthesis, which in that case could occur during the proliferation of erythrocytes. In a separate study we found that such an induction does take place in smokers as GST concentrations determined by ELISA increased, whereas the activity towards CDNB was not increased [7]. In the current exposure situation a decrease in activity due to inactivation of GST is expected to prevail over induction because the duration of the exposure period was short compared to the life span of an erythrocyte. Normal GST activity shows appreciable variation between subjects [21]. In this respect, the comparison of individual pre- and post-exposure values is expected to be more sensitive than comparison with a control group. Until now, no genetic inhomogeneity has been found to explain the interindividual variations in erythrocyte GST activity towards CDNB (mainly GST activity). Such genetic differences are known to exist for GST μ [25] and are expected for the minor erythrocyte GST [19]. Therefore, the absence of a relation between erythrocyte GST and the presence or absence of GST μ in lymphocytes was not surprising. The distinct decrease of both GSH and GST values found in this study indicates that exposure to DCP has affected glutathione conjugating capacity in erythrocytes.

Acknowledgments. We wish to thank A.J. Moolenaar, J.H.M. Souverijn, M.A. Verschoor, A.M. van Schijndel, H. de Jour and G.J. Reijmer for their analytical support, and D.H. Brouwer, J.J. van Hemmen, R.P. Bos, P.Th. Henderson, and R.L. Zielhuis for their valuable comments on the manuscript. This study was financially supported by the Food and Allied Workers Union.

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Chapter 8

Changes in Blood Glutathione Concentrations, and in Erythrocyte Glutathione Reductase and Glutathione S-Transferase Activity After Running Training and After Participation in Contests

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Published in: *Eur J Appl Physiol* **64**, 1992, 354–358.

Summary. Previously sedentary men ($n = 23$) and women ($n = 18$) were trained to run a half marathon contest after 40 weeks. Total blood glutathione had increased by 20 weeks of training and had returned to normal after 40 weeks. Erythrocyte glutathione reductase activity had increased by 20 weeks and remained elevated after 40 weeks. This effect was accompanied by decreases in glutathione reductase coefficients, which indicated that increases in the presence of riboflavin may have been responsible for the changes in reductase activity. Erythrocyte glutathione S-transferase activity had increased slightly after 20 weeks of training and a much more marked increase was found after 40 weeks. This may have been indicative of the occurrence of lipid peroxidation in this phase of training. The participants ran a 15-km race after the first 20 weeks of training and a half marathon after 40 weeks. Blood glutathione tended to decrease after the 15-km race and increased after the half marathon. In both cases it had returned to normal values 5 days after the race. Erythrocyte glutathione reductase was elevated 1 day after the races, and had returned to normal after 5 days. This could also have been explained from concurrent changes in the riboflavin content of the erythrocytes. Erythrocyte glutathione S-transferase activity decreased after both races, but was restored 5 days after the half marathon while such a restoration was not the case after the 15-km race.

Introduction

Endurance training has been shown to have positive effects with respect to endurance capacity, metabolic stress tolerance and a number of haematological parameters [8]. Increased physical activity leads to increased oxygen consumption. Since 1%–2% of the oxygen consumed gives rise to reactive oxygen species, this also has led to an increased challenge to the antioxidant system[9]. Part of this change in the antioxidant system may also result from the partial ischaemia in active tissues that may have occurred during heavy exercise[2], or from the release of heat shock proteins in these tissues[6]. Lipid peroxidation initiated by free radical reactions has occurred in muscle after heavy exercise, and gave rise to increased thiobarbituric acid reactive substances which were accompanied by increased activities of creatine kinase, lactate dehydrogenase and aspartate aminotransferase in serum[14, 15]. Especially in untrained rats increased levels of tissue lipid peroxidation have been accompanied by elevated hepatic catalase and cytosolic superoxide dismutase activities [10]. Increased oxygen turnover could also lead to an increased formation of reactive oxygen species in erythrocytes. Increased methaemoglobin formation alone would already provide a challenge for glutathione-dependent reducing systems. Ohno *et al.* [16] have reported increases in erythrocyte glutathione reductase (GR) activity and in GR coefficients in seven males after 10 weeks of training. Indications of radical stress in blood have also

been found by Pincemail *et al.* [17], who showed that, after exercise on a cycle ergometer increased concentrations of the radical scavenging vitamin tocopherol were found not only in plasma but also in the erythrocyte.

To gain a better insight of the exact nature of the effects of exercise on the glutathione dependent defense system in the erythrocyte, we studied the effects of running training on total blood glutathione (GSH), on GR and glutathione S-transferase (GST) activities in erythrocytes and on the riboflavin content of the erythrocytes. Blood samples were collected before the start of the experiment, after 20 weeks of training and after 40 weeks of training. The 20- and 40 week samples were collected before the subjects studied participated in a 15-km and a half marathon contest, respectively. The effects of the contest itself were studied in samples taken before and after the races.

Methods

Study design

The participants (23 men and 18 women) were selected from a population of 370 who replied to a bulletin issued through the local media. Prior to the study they had not participated in any running sport and were not active in any other sport for more than 1 h a week. The selection was made to produce a comparable group of both sexes in respect of age (28–41 years) and body mass index (19.4–26.4 kg·m⁻²). The subjects were examined medically and gave their written informed consent before the start of the study.

The training programme has been described before [8]. In brief, it consisted of three to four prescribed training sessions weekly, one of which was supervised by one of the authors (GJ). The time spent running was gradually increased from 10 to 30 min during the first 8 weeks, to 20 to 60 min in weeks 9 to 20 and to 30 to 90 min in the last part of the study.

Two participants of each sex did not continue the training until the first 20 weeks, and were removed from the study. Due to personal circumstances no samples were collected from 4 women and 1 men before and after the 15-km race. One woman and 3 men abandoned the study during the second period of training and, therefore, did not participate in the half marathon contest.

To study the effect of training, blood samples were collected before the start of the experiment, after 20 weeks of training, and after 40 weeks of training. To evaluate the effects of participation in contests blood samples were taken 5 days before (day -5), immediately after (day 1) and 5 days after (day 5) a 15-km contest ran after 20 weeks and a half marathon run after 40 weeks.

The GSH concentrations in blood, GR [Enzyme Commission no. (EC) 1.6.4.2] and GST (EC 2.5.1.18) activities in erythrocytes and the glutathione reductase coefficient (GR_{coeff.}, an indicator of the riboflavin content) of the erythrocytes were determined in all samples.

Analytical methods

Chemicals.

The following chemicals were used: 5,5'-dithiobis-(2-nitrobenzoic acid) (*DTNB*; E. Merck, Darmstadt, FRG). Oxidized (*GSSG*) and reduced (*GSH*) glutathione (Boehringer Mannheim, Mannheim, FRG), 1-chloro-2,4-dinitrobenzene (*CDNB*), flavin adenine dinucleotide (*FAD*), nicotinamide adenine dinucleotide phosphate, reduced (*NADPH*) and GR (Sigma; St. Louis, MO, USA). All other chemicals were of analytical quality. Only microfiltrated deionized water was used.

Determination of blood GSH.

Blood samples for determination of total GSH were collected in 5-ml vacutainers containing 7.5 mg Na_2EDTA (ethylenediaminetetraacetic acid). To 1-ml blood an equal volume of 14% (v/v) HClO_4 was added. After centrifugation at 3000 *g* for 10 min the supernatant was neutralized with a mixture of 0.3 mol·l⁻¹ N-2-hydroxyethylpiperazine-N'-2-ethanesulfonic acid (buffer) (HEPES) and 2 mol·l⁻¹ KOH. The samples were stored below -70°C until determination. Total GSH was determined with the cyclic oxidation-reduction method essentially as described by Anderson [1].

Determination of enzyme activities.

Erythrocytes isolated from 10-ml blood samples collected in vacutainers containing heparin, were washed three times with saline and stored below -70°C until further use. To the frozen erythrocyte samples 1-ml water was added, and the samples were thawed on ice. Thereafter, the samples were diluted twenty fold with 100 mmol·l⁻¹ potassium phosphate buffer (pH=6.5) containing 6.3 mM Na_2EDTA and centrifuged at 10.000 *g* for 10 min. The GST activity towards CDNB was determined in 200 μl of this sample as described by Habig and Jakoby [7] with a Philips PU 8740 spectrophotometer (Philips; Eindhoven; NL) with the slit adjusted to 2 nm. A quantity of 200 μl of the same sample was used for determination of GR activity according to the method described by Carlberg and Mannervik [4]. The haemoglobin concentrations in the erythrocytes were determined with the haemoglobin cyanide procedure described by Van Kampen and Zijlstra [12].

Determination of riboflavin content.

The coefficient of erythrocyte GR activity with and without addition of FAD to the assay was used as an indicator of the riboflavin content. The determinations were carried out essentially as described for GR above, except that 0.01 mM FAD was added to one of the two samples[18].

Statistical analyses.

Sex related differences were evaluated with the Mann-Whitney *U*-test. Differences between values on different days were analysed with Wilcoxon's matched pair analysis with matches for values from the same individual.

Results

Sex differences.

Sex related differences were evaluated for data combined for all days. The GSH values were found to be higher in men than in women [1.14 (SD 0.34) mM and 1.06 (SD 0.35) mM, respectively; $p < 0.029$]. This difference resulted directly from the difference in packed cell volume between men and women, and disappeared when GSH concentrations were expressed as mmol·g Hb⁻¹. The GR activity was higher ($p < 0.005$) and the GR_{coeff.} were lower ($p < 0.046$) in women. The values were 3.29 (SD 0.56) U·g Hb⁻¹ with a ratio of 1.05 (SD 0.09) for men and 3.5 (SD 1.2) U·g Hb⁻¹ with a ratio of 1.03 (SD 0.10) for women. Since the statistical procedures were carried out as matched pair analyses, and since there were no qualitative differences between sexes in the changes found, all reported changes were calculated for men and women taken together. There was one exception, and that was that the glutathione concentration increase after the 1/2 marathon (see below) did only occur in females, while in males an opposite (not statistically significant) tendency existed.

Values after training.

Changes in blood GSH and erythrocyte GST, GR and GR_{coeff.} values during the training period are given in Table 8.1. Total GSH in blood was increased almost 50% after the first 20 weeks of training. After the prolonged (heavy) training in the second part of the study, the values returned to the initial levels and were in fact even slightly lower. The GR activity had also increased [from 3.13 (SD 0.76) U·g Hb⁻¹ to 3.36 (SD 0.85) U·g Hb⁻¹] after the first 20 weeks and was still at a high level [3.42 (SD 1.01) U·g Hb⁻¹] after 40 weeks. Accompanying this change in GR activity, we found a decrease in the GR_{coeff.} values [from 1.08 (SD 0.09) to 1.04 (SD 0.03)] in the first 20 weeks. Like the GR values these values showed no further changes during the second training period. The GST activity increased only slightly during the first period [1.08 (SD 0.41) U·g Hb⁻¹ before and 1.15 (SD 0.43) U·g Hb⁻¹ after 20 weeks] but increased markedly during the second part of the study [to 1.86 (SD 0.75) U·g Hb⁻¹].

Participation in contest.

Changes in blood GSH and erythrocyte GST, GR and GR_{coeff.} values after participation in the 15-km race and in the half marathon and the values for all four

Table 8.1: Effects of training on glutathione concentrations (GSH) in blood, glutathione S-transferase (GST) and glutathione reductase (GR) activity and the glutathione reductase activity coefficient ($GR_{\text{coeff.}}$) in erythrocytes. The training schedule lasted 40 weeks and for the values after 20 weeks and 40 weeks levels of significance for differences with the earlier samples (Wilcoxon matched pair test) are given. p Values > 0.1 are given as NS (not significant).

			level of significance for difference from:	
			0 weeks	20 weeks
GSH	0 weeks	6.54 (SD 2.26) mmol·g Hb ⁻¹		
	20 weeks	9.73 (SD 2.09) mmol·g Hb ⁻¹	$p < 0.0001$	
	40 weeks	5.94 (SD 1.61) mmol·g Hb ⁻¹	NS	$p < 0.0001$
GR	0 weeks	3.13 (SD 0.76) U·g Hb ⁻¹		
	20 weeks	3.36 (SD 0.85) U·g Hb ⁻¹	$p < 0.0037$	
	40 weeks	3.42 (SD 1.01) U·g Hb ⁻¹	$p < 0.0003$	NS
$GR_{\text{coeff.}}$	0 weeks	1.08 (SD 0.09)		
	20 weeks	1.04 (SD 0.03)	$p < 0.015$	
	40 weeks	1.03 (SD 0.10)	$p < 0.016$	NS
GST	0 weeks	1.08 (SD 0.41) U·g Hb ⁻¹		
	20 weeks	1.15 (SD 0.43) U·g Hb ⁻¹	$p < 0.018$	
	40 weeks	1.86 (SD 0.75) U·g Hb ⁻¹	$p < 0.0001$	$p < 0.0001$

parameters determined 5 days after each race are given in table 8.2. For both races the GSH values in the samples taken 5 days after the race were not statistically different from the precontest values. The GSH concentrations tended to be lower after the 15-km race. While the decrease from pre- to postrunning samples did not attain statistical significance, the p value for the subsequent increase (equal to a return to normal values) was 0.066. Since the precontest values were in fact increased in comparison with the samples taken at the start of the project, the decrease went towards normal values. After the half marathon we found an increase in GSH values. As the precontest values for the half marathon were lowered in comparison with the pretraining values, here too the shift was in the direction of the pretraining values.

The GR activities were increased significantly ($p < 0.03$) in the day 1 samples taken after both contests and had returned to prerunning values on day 5. Here too, the change was accompanied by a change in $GR_{\text{coeff.}}$ values. There was a slight decrease in these values after both contests (not significant after the half marathon) and a return to normal at day 5 (the increase between

Table 8.2: Effects of participation in a 15-km and a half marathon contest on glutathione concentrations (GSH) in blood, glutathione S-transferase (GST) and glutathione reductase (GR) activity and the glutathione reductase activity coefficient (GR_{coeff.}) in erythrocytes. Values are given for the participants 5 days before (-5), 1 day after (+1) and 5 days (+5) after the contests. For the values at days 1 and 5 levels of significance for differences with the earlier samples (Wilcoxon matched pair test) are given. *p* Values > 0.1 are given as NS (not significant).

			level of significance for difference from values at: 5 days before 1 day after race (-5) race (+1)	
GSH	15km -5	9.73 (SD 2.09) mmol·g Hb ⁻¹		
	15km +1	9.34 (SD 1.62) mmol·g Hb ⁻¹	NS	
	15km +5	9.77 (SD 1.85) mmol·g Hb ⁻¹	NS	<i>p</i> < 0.066
	$\frac{1}{2}$ mar -5	5.94 (SD 1.61) mmol·g Hb ⁻¹		
	$\frac{1}{2}$ mar +1	6.52 (SD 1.51) mmol·g Hb ⁻¹	<i>p</i> < 0.057	
	$\frac{1}{2}$ mar +5	6.11 (SD 1.83) mmol·g Hb ⁻¹	NS	NS
GR	15km -5	3.36 (SD 0.85) U·g Hb ⁻¹		
	15km +1	3.47 (SD 0.91) U·g Hb ⁻¹	<i>p</i> < 0.021	
	15km +5	3.35 (SD 0.87) U·g Hb ⁻¹	NS	<i>p</i> < 0.0067
	$\frac{1}{2}$ mar -5	3.42 (SD 1.01) U·g Hb ⁻¹		
	$\frac{1}{2}$ mar +1	3.56 (SD 1.02) U·g Hb ⁻¹	<i>p</i> < 0.027	
	$\frac{1}{2}$ mar +5	3.44 (SD 1.04) U·g Hb ⁻¹	NS	<i>p</i> < 0.032
GR _{coeff.}	15km -5	1.04 (SD 0.09)		
	15km +1	1.02 (SD 0.09)	<i>p</i> < 0.010	
	15km +5	1.03 (SD 0.09)	NS	NS
	$\frac{1}{2}$ mar -5	1.03 (SD 0.10)		
	$\frac{1}{2}$ mar +1	1.02 (SD 0.11)	NS	
	$\frac{1}{2}$ mar +5	1.04 (SD 0.11)	NS	<i>p</i> < 0.022
GST	15km -5	1.15 (SD 0.43) U·g Hb ⁻¹		
	15km +1	1.12 (SD 0.40) U·g Hb ⁻¹	<i>p</i> < 0.077	
	15km +5	1.10 (SD 0.41) U·g Hb ⁻¹	<i>p</i> < 0.018	<i>p</i> < 0.081
	$\frac{1}{2}$ mar -5	1.86 (SD 0.75) U·g Hb ⁻¹		
	$\frac{1}{2}$ mar +1	1.64 (SD 0.60) U·g Hb ⁻¹	<i>p</i> < 0.0022	
	$\frac{1}{2}$ mar +5	1.91 (SD 0.73) U·g Hb ⁻¹	NS	<i>p</i> < 0.0001

day 1 and day 5 was significant for the half marathon).

The GST activities had decreased after both the 15 km contest and the half marathon. This change was much more pronounced for the half marathon samples but it should be noted that the precontest values for this race were increased compared with the pretraining values. In the day 5 samples taken after the 15-km contest the GST activity values were even further decreased, whereas the values at day 5 after the half marathon had returned to precontest values.

Discussion

We found increases of the order of 50% for GSH after 20 weeks of training. An increase in blood GSH during a period of training can be explained by the high amounts of reduced GSH that are needed for the reduction of methaemoglobin formation and for protection against reactive oxygen species. The same argument applies to the (less pronounced) increase in GR activity. During the second part of the study the GSH values returned to normal. This phenomenon could be explained by a shortage in glutathione synthesis or by the disappearance of the need for elevated glutathione concentrations. Increases in GR activity could be a possible cause for such a diminution in the need for GSH availability, as higher GR activity levels could have resulted in a shift of the ratio of GSH and GSSG. The continuously high levels of GR activity found in the second part of the training period, indicated that a decrease in the need for GSH was not likely. The increase in GR activity was, at least in part a result of increases in riboflavin availability, which was indicated by decreased $GR_{\text{coeff.}}$ values in the samples collected after the training periods. The findings of Ohno *et al.*[16] would suggest that this was not the only cause. While they have also reported increases in GR activity after training they have found increased $GR_{\text{coeff.}}$. The differences in the procedures used in these two studies may have accounted for this difference in the effects on $GR_{\text{coeff.}}$ (only 7 subjects and only 10 weeks of training were used by Ohno *et al.* 1988). Training would seem to increase the need of erythrocytes for reduced GSH. While this need was initially countered by increased GSH synthesis, later on only increases in GR activity were found. The presence of relatively large amounts of younger cells in the 20 week samples may also have contributed to the explanation of the increased GSH concentrations. This was indicated by the lower ($p < 0.0001$) mean cell volumes (MCV) in these samples [MCV values were 93.2 (SD 4.1) fl in the pretraining period; 91.5 (SD 4.1) fl in the 20-week and 92.6 (SD 3.8) in the 40-week samples]. This decrease in MCV in the early training phase was in accordance with what has been found in a previous study [11]. While there was only a small increase in the level of GST activity during the first 20 weeks of training, a marked increase occurred during the second period. Since GST is vulnerable in the presence of reactive species (see below), which could be seen in the decrease of GST activity after the races, this increase may have been a result

of a kind of overshoot adaptation to such repeated inactivations. On the other hand, GST apart from its functions as a transferase and an intracellular transport protein, has an important function as an organic peroxidase [13]. Exercise can lead to increased lipid peroxidation in muscles [14,15]: our findings indicated that lipid peroxidation after exercise may also have occurred in erythrocytes.

In previous studies [3] a decrease in GST activities in erythrocytes after exposure to the soil fumigant dichloropropene has been reported. The decrease found after heavy physical activity may also have been indicative of the formation of reactive substances. The return to normal values 5 days after the half marathon (but not after the earlier 15-km race) indicated that GST inactivation was reversible and suggested that the reactive species were either formed less or removed faster in longer trained persons. The increase of GR activity directly after the races could hardly have been a result of *de novo* synthesis of GR, since erythrocytes lack active protein synthesis, and since red-cell proliferation is slow. The concurrent changes in GR activity coefficients indicated that the changes in GR activity were most likely to have been caused by modulations in activity as a result of changes in erythrocyte riboflavin concentrations. The changes in GSH after the races were minor. Duthie *et al.* [5] have reported significant ($\pm 15\%$) decreases of reduced GSH in samples taken 24 h after a half marathon. Both the direction and the magnitude of this change are not in agreement with the values reported here.

In general it can be concluded that both endurance training and competitive physical exercise had important effects on the glutathione dependent defense system in erythrocytes. Most effects appeared to be reversible adaptations. The pronounced increase in GST activity during the second phase of the training programme may, however, indicate a response to cell damaging reactions.

Acknowledgment. We thank A.J.P. van Bommel for his technical assistance.

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Chapter 9

Decreased Glutathione Content and Glutathione S-Transferase Activity in Red Blood Cells of Coal Miners with Early Stages of Pneumoconiosis

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Published in: *Br J Ind Med* 50, 1993, 633–636.

Summary. Blood samples of miners heavily exposed to coal dust were examined for changes in glutathione S-transferase (*GST*). Decreased *GST* activity was found in red blood cells of subjects at early stages of coal workers' pneumoconiosis (International Labour Office classification 0/1–1/2) when compared to control miners. At further progression of coal workers' pneumoconiosis ($\geq 2/1$), the activity of *GST* was not different from controls. In the same group with moderate coal workers' pneumoconiosis a decrease in GSH in red blood cells occurred. Decreases in *GST* activity in early stages of coal workers' pneumoconiosis, as well as the decreases in glutathione peroxidase (*GPX*) activity and in GSH reported earlier [9], may originate from damage caused by reactive oxygen species. These changes might imply an impairment of the detoxification capacity for electrophilic and oxidative compounds during this stage of the disease.

Introduction

Lung fibrosis related to the inhalation of dusts containing toxic particles such as silica, or asbestos is a serious occupational hazard. It is well known that during the immune defense against dust particles, reactive oxidant species are secreted by various types of inflammatory cells with the alveolar macrophage playing a key role in this process [5, 21]. The lung tissue possesses an elaborate defense mechanism to detoxify reactive oxygen species, constituted by a set of antioxidant enzymes (*AOE*) such as superoxide dismutase and glutathione-peroxidase (*GPX*) and low-molecular compounds such as glutathione (*GSH*) acting as a substrate for *GPX* and glutathione S-transferase (*GST*) or as direct radical scavengers. Also, antioxidants localised in erythrocytes are involved in the antioxidant protection of the lung [2, 22], acting as circulating antioxidant carriers.

Previously, we showed [9, 3, 4] that *AOE* and *GSH* in erythrocytes are affected at different stages of pneumoconiosis in coal workers or silicotic patients and in lung tissue of rats exposed to silica [13]; *GSH* concentration and *GST* activity were also found to be affected after occupational exposure to reactive compounds [6, 10] and in situations involving reactive oxygen species stress like smoking [10] and heavy exercise [11]. This paper presents the results of *GSH* determinations and *GST*-analyses with an activity assay and enzyme linked immunosorbent assay (*ELISA*) in red blood cells of a cohort of coal miners previously investigated by our research group [9].

Materials and Methods

Subjects

Ninety one coal miners, all heavily exposed to coal dust underground at the coal face for at least twelve years, were selected from the Belgian coal mining industry

pits (Kempense Steenkolenmijnen). Blood samples were taken and treated as described previously [9]. Control miners ($n = 58$) and miners with coal workers pneumoconiosis (*CWP*) ($n = 33$) were matched for exposure and age at which they started work. In this study persons with the classification 0/1, 1/0, 1/1 and 1/2 [17] were gathered in group 2 ($n = 19$); persons with classification 2/1 and higher in group 3 ($n = 14$); control miners constituted group 1.

Analytical methods

Chemicals.

The chemicals used were: rabbit polyclonal antibodies against GST π (Medlabs; Dublin, Ireland), swine anti-rabbit IgG antibodies conjugated to horse-radish peroxidase (*HRP*) and orthophenylenediamine (*OPD*) (Dakopatts; Copenhagen, Denmark), 5,5'-dithio-bis(2-nitrobenzoic acid) (*DTNB*); E. Merck; Darmstadt, FRG). Reduced glutathione (*GSH*) (Boehringer Mannheim; Mannheim, FRG), 1-chloro-2,4-dinitrobenzene (*CDNB*), dithiothreitol (*DTT*), GST π from human placenta and tween-20 (Sigma; St. Louis, MO, USA). All other chemicals were of analytical quality. Only microfiltrated deionized water was used.

Determination of blood GSH.

Total GSH was determined with the cyclic oxidation-reduction method essentially as described by Anderson [1].

Assays for GST.

GST (EC 2.5.1.18) activity was determined by a modification of the method described by Habig and Jakoby [12] as previously described [11] lysing the RBC by addition of three volumes of water containing 1.4 mM neutral DTT, centrifugation and measuring the activity towards CDNB. A Cary 118 spectrophotometer was used in auto slit mode with the gain adjusted to 2.5.

For the competitive ELISA, CostarTM flat bottom 96-well EIA plates (Data Packaging Corporation; Broadway, MA, USA) were coated overnight at 4°C with 200 μ l phosphate buffered (10 mM) saline of pH 7.2 containing 11 μ U GST π from human placenta per well. Each well was washed three times with 200 μ l ELISA buffer (10 mM sodiumphosphate + 0.5 M sodiumchloride + 0.1% (v/v) Tween-20) of pH 7.2. This and all further steps were performed at room temperature. To each well 50 μ l of the antigen solution (hemolysate) and 150 μ l antibodies against GST π diluted 2250 fold with ELISA buffer were added. After 2 hours incubation the wells were washed three times with ELISA buffer and incubated for 1 hour with 200 μ l HRP labeled anti-IgG diluted 1500 fold with ELISA buffer. After another three washes with ELISA buffer 100 μ l of a 100 mM citrate-phosphate buffer of pH 5.0 containing 0.67 mg/ml OPD and 0.0125 % (w/v) H₂O₂ were added to each well. After exactly 15 minutes the enzymatic reaction was stopped

by addition of 150 μ l 1 *M* sulfuric acid. The extinctions at 492 nm were read with a multiscan interfaced to a MINC computer.

Reference data for the quantitation of the ELISA results were obtained with GST π from human placenta. Known quantities of this enzyme —expressed in activity units— were tested in the competitive ELISA assay. The results were fitted to the general logistic function [7]:

$$Y = \frac{a - d}{1 + \left(\frac{C}{EC_{50}}\right)^s} + d$$

where Y is the response; C the concentration of the antigen; a the response when $C = 0$; d the response for “infinite” concentrations of antigen; EC_{50} the antigen concentration resulting in a response halfway between a and d ; and s is a slope factor which corresponds to the slope of the logit-log plot, when C is plotted in terms of natural logarithms. For curve fitting the NLIN procedure from the SASTM package was used with the DUD [20] method for nonlinear regression. The response was expressed as fraction of the maximum extinction, found without addition of antigen, $E = \frac{Y}{a}$.

The haemoglobin concentrations in the erythrocytes were determined with the haemoglobin cyanide procedure [14].

Statistical Evaluation

Analytical results were coupled to previous data on this cohort in a database. Correlations between variables were studied using STATGRAPH 3.01 statistical package. Differences between variables in separate groups were evaluated with the Wilcoxon 2 sample test.

Results

Table 9.1 shows the mean values and standard deviations (SDs) of GSH concentrations, GST activity towards CDNB and GST content as determined by competitive ELISA in the blood of the control miners and the two groups of miners with coal workers' pneumoconiosis. No effect of smoking was seen on any parameter mentioned. GSH values were determined previously [9]. Repeated analyses in the frozen samples used for the GST determinations did not show significant differences from the original values. Therefore, the original data rearranged for the current purpose are listed in table 9.1. Both GSH concentration and GST activity values were decreased in group 2, the workers with moderate forms of coal workers' pneumoconiosis. In group 3 such a decrease was not found. The GST protein content showed a tendency to follow the GST activities, although the decrease in group 2 did not attain statistical significance.

Table 9.1: Glutathione (GSH) concentrations, glutathione S-transferase (GST) activity towards CDNB and glutathione S-transferase protein content as determined by ELISA in red blood cells of coal workers. Levels of significance for decrease compared to the control group (Wilcoxon 2 sample test) are given for *p* values smaller than 0.1.

	Control miners (0/0)	Group 2 CWP (0/1-1/2)	Group 3 CWP ($\geq 2/1$)
GSH (mmol·g Hb ⁻¹)	3.97 (SD 0.08) (<i>n</i> = 58)	3.62 (SD 0.14) (<i>n</i> = 19; <i>p</i> = 0.016)	3.98 (SD 0.46) (<i>n</i> = 14)
GST (CDNB) (U·g Hb ⁻¹)	2.99 (SD 1.36) (<i>n</i> = 45)	2.31 (SD 1.09) (<i>n</i> = 9; <i>p</i> = 0.061)	3.37 (SD 1.14) (<i>n</i> = 10)
GST (ELISA) (\equiv U·g Hb ⁻¹)	2.46 (SD 1.93) (<i>n</i> = 34)	1.92 (SD 0.90) (<i>n</i> = 6)	2.54 (SD 1.39) (<i>n</i> = 7)

Discussion

Our data show that, as well as GSH, erythrocyte GST activity is decreased in subjects with early stages of pulmonary fibrosis (0/1-1/2) when compared with control miners. At further progression of coal workers' pneumoconiosis (2/1 +) however, erythrocyte activity was not different from controls. In the same group with moderate coal workers' pneumoconiosis, a decrease in GSH in erythrocytes occurs. Previously, we reported an increase in total GPX activity in the same workers at early stages of the disease [9] and in rat lung homogenates after *in vivo* exposure to silica or asbestos [13]. We think that the decrease in GSH and GST activity is caused by excessive release of reactive oxygen species by alveolar macrophages and neutrophils in the lung tissue. A decrease in GST activity cannot be the result of decreased GSH concentrations during determinations, since saturating concentrations of GSH (1 mM) are added in the assay. Moreover, there is no (negative) correlation between GSH content and GST activity of the blood samples. Combined with our data obtained with the ELISA on GST in erythrocytes (which tended to decreased in the same group) we conclude that a fraction of GST is damaged, possibly by oxidation of SH-sites. The decrease in available GST activity plus the fact that GST activity is GSH dependent, might imply that *in vivo* functioning of the enzyme is affected rendering these individuals more susceptible to damage resulting from lipid peroxidation or from co-exposure to electrophilic compounds. Various mineral fibers were reported [19] to inhibit GST activity in cytosol of rat lung homogenates after *in vitro* and *in vivo* exposure, while phase 1 reactions were increased at advanced stages of the

disease. An induction of phase I enzymes in advanced stages of fibrosis might further aggravate the accumulation of activated metabolites in the lung.

Previously, we argued that the restoration of erythrocyte GSH content in miners with coal workers' pneumoconiosis beyond 2/1 to control values, is caused by a hepatic efflux of GSH [9]. We found comparable phenomena for GST in sporters. After moderate running training (up to 60 min after 20 weeks) [11] GST activity and GSH concentrations in RBC were both increased. After heavy training (up to 90 min after 40 weeks) GSH values were returned to normal and GST was strongly increased. Increases in GST protein content in RBC can be the result of increased protein synthesis during RBC proliferation, or, less likely, of increased longevity of the protein itself. Increases in GST content were also found in smokers [10]. Here lower specific activities and equal total activities were found. This too might be a result of compensatory GST synthesis. Thus, GST activity in coal miners with coal workers' pneumoconiosis (>2/1) could remain at normal values due to compensation of inactivation by extra protein synthesis during RBC proliferation. This extra synthesis could be triggered by inflammatory processes generating cytokine release in coal workers' pneumoconiosis[15] or very heavy exercise. Possibly the formation of lipid peroxides, documented in exercise [8] and rats exposed to mineral fibers[18], could be an important factor in this respect. Lipid peroxides are known to be detoxified by GST, which shows selenium independent GPX activity [16].

In conclusion, red blood cell GST activity is affected in early phases of coal workers' pneumoconiosis, possibly as a result of oxidative damage. Combined with our earlier findings (a decrease in GSH, and increase of GPX) this might mean that detoxification of electrophilic compounds in this stages is impaired.

Acknowledgement. The authors wish to thank drs. K. Herberichs and J.G.P. Peters of the Department of toxicology from the University of Nijmegen for their assistance with the analyses.

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Chapter 10

General Discussion

The presence of reactive chemicals within the human body may lead to a disruption of the integrity of various biological macromolecules including nucleic acids. Toxicological stress factors that produce such effects can roughly be divided in two types: electrophilic compounds and oxidative stress. Exposure to a broad variety of chemicals can give rise to increased electrophilic stress, either because of the direct reactivity of these xenobiotics or because of the formation of reactive metabolites. When reactive electrophilic compounds reach nucleophilic centers in biological macromolecules like DNA or protein, they can react directly, often leading to the formation of covalent bindings. The exact consequences of such reactions depend on the identity of the biological target and the nature and stability of the adduct. At low exposures to electrophiles, such as may occur during occupational exposure to chemicals used at the workplace or for instance as a result of cigarette smoking, the occurrence of mutations in the nucleic acid sequence is the most important risk factor. Oxidative stress can originate from a broad gamut of conditions ranging from endogenous formation of active oxygen species during normal metabolism and inflammatory activity [26] to radical formation upon exposure to oxidative compounds. As a result of oxidative stress, free radical reactions may occur which can cause damage to nucleic acid and protein structures. The most prominent target for such radical reactions however, are the lipid structures in biological membranes. Especially polyunsaturated lipids tend to propagate radical chain reactions. During this lipid peroxidation process reactive products like aldehydes and lipid peroxides are formed, which can cause secondary damage to other biological structures.

The glutathione – glutathione S-transferase system described in this thesis is involved both in the protection against electrophiles and in the reduction of oxidative stress. The glutathione S-transferase (*GST*) catalysed phase II conjugation of electrophilic compounds with reduced glutathione (*GSH*) is the important first step in the formation of mercapturic acid excretion products [32]. Usually this results in detoxification. *GSH* itself is consumed in the transferase reactions, and since the thioethers formed are removed from the cell [2] this may lead to a depletion of *GSH*. Furthermore, *GST* may also become a target for electro-

philic compounds. This suicidal reaction leads to loss in GST activity [8,20,39]. This phenomenon also occurs when human red blood cells are exposed *in vitro* to important industrially used compounds like acrolein, propylene oxide, styrene oxide, ethylene dibromide and ethylene dichloride [1].

The involvement of GSH in protection against oxidative stress is of a more complex nature [5,29]. GSH can be used to detoxify hydroperoxides generated as a result of oxidative stress (equation 2.1; p. 18). Selenium dependent glutathione peroxidases catalyse this reaction for both hydrogenperoxide and lipidperoxides. Scavenging of hydrogenperoxide is of particular interest since it can react with superoxide ($O_2^{\cdot-}$) radicals, thereby generating the potent hydroxyl (HO^{\cdot}) radical. Removal of lipidperoxides is of importance since these reactive compounds can damage other biological structures and because they are potentially involved in amplification of radical reactions [eqs. 2.18 and 2.19; p. 26]. For lipidperoxides the peroxidase reaction is also catalysed by the nonselenium GST. GSH might also protect by direct free radical scavenging, but the importance of this mechanism is not clear. The effectiveness of such direct scavenging is further limited by the presence of GSH in the aqueous phase, which makes it impossible to terminate lipid radical propagation. GSH may however have another important contribution to free radical scavenging. The main lipid soluble antioxidant is α -tocopherol (vitamin E). GSH is expected to be able to reduce the tocopheroxyl radical formed from radical scavenging by α -tocopherol [18,24]. Apart from this, ascorbate (vitamin C) is well known for the same kind of tocopherol restoring activity [4,24]. Therefore, GSH might also aid in the restoration of α -tocopherol indirectly by reducing dehydroascorbate. A glutathione dehydroascorbate oxidoreductase is present in plant tissues. It was shown recently that dehydroascorbate reduction by GSH is catalysed by thioltransferases from porcine liver, human placenta and bovine thymus as well as by bovine protein disulfide isomerase [41]. This lead Wells *et al.* to the suggestion that the same activity might be present in erythrocytes [42]. They did not actually demonstrate the activity in human erythrocytes however. The findings by Stahl and coworkers, who took a closer look at the glutathione dehydroascorbate reductase activity in human leucocytes and only found evidence for nonenzymatic activity [33], show how important such an actual demonstration is. GSH also functions as a general buffer for redox equivalents. This is possible because of the high concentration of GSH in red blood cells (about 2 mM) and because of the normally very high redox potential of the GSH/GSSG couple. Oxidation of GSH will result in an activation of glutathion reductase and in a GSSG and glutathione reductase mediated activation of glucose 6-phosphate dehydrogenase [19]. As long as glucose supply is sufficient this will result in a rapid recovery of homeostatic GSH/GSSG ratios. Another important function of GSH is the protection of protein thiol groups. GSH can reduce disulfides and sulfoxides formed as a result of oxidative reactions thus restoring damaged and thereby often inactivated proteins [9,21,22]. Following oxidative stress conditions mixed protein-glutathion disulfides are of

ten detectable. Since oxidized glutathione *GSSG* is toxic to erythrocytes, storage of oxidized glutathione as mixed disulfides might be more beneficial. In fact the quantitative importance of this mixed disulfide formation is larger than *GSSG* excretion, at least under some *in vitro* conditions studied (Chapter 5 and [12]). Since all protective effects described will lead either to the excretion of *GSSG* from the erythrocyte or to the fixation of free glutathione as mixed disulfides, the net result of oxidative stress will also be a decrease in erythrocyte *GSH*. The analogy between the effects of the two stress factors goes even further. The main glutathione *S*-transferase isoenzyme present in human erythrocytes belongs to the π class [11]. In studies published during the last few years evidence was presented that this class of *GST* is vulnerable to oxidative stress [30,31,37]. Our own *in vitro* experiments (not shown in this thesis) confirmed that human erythrocyte *GST* activity can also be strongly impaired by hydrogenperoxide treatment.

From the above it can be concluded that it is conceivable that exposure to electrophilic or oxidative compounds, or to any other kind of oxidative stress, would lead to decreases in glutathione content or glutathione S-transferase activity in erythrocytes. The studies described in this thesis were focussed on the practical usefulness of this idea for experimental toxicological research and for biological effect monitoring purposes. The first part (chapter 3-5) consists of in vitro studies using human blood. In the second part (chapter 6-9) a number of in vivo biological effect monitoring studies are described.

Directly reactive electrophiles like iodoacetamide, diethyl maleate and *N*-ethylmaleimide have a high reactivity towards glutathione. All these compounds gave strong *GSH* depletions both in hemolysates and in intact erythrocytes (Chapter 3). As a result of this, *GSH* is able to protect other molecules present in the erythrocytes. The formation of hemoglobin adducts showed a strong increase when reagent concentrations above that leading to full *GSH* depletion were used. Modulation of the *GSH* related protection *in vivo* will also have consequences for the formation of hemoglobin adducts. This may be of practical importance since measurement of hemoglobin adducts is itself used as a biological effect monitoring parameter for exposure to electrophilic genotoxic chemicals [10,23,38]. Especially the development of nonlinear models for hemoglobin adduct formation [6] will benefit from increased understanding of this kind of protection mechanisms.

The cytostatic drug cisplatin is known to covalently bind towards DNA and proteins [7]. It also caused depletion of total (=reduced + oxidized) glutathione (*GT*) in isolated cytosol of human red blood cells and in rat hepatocytes or rat liver cytosol (chapter 4). In intact human red blood cells no *GT* depletion was found, probably because the erythrocyte membrane effectively prohibits uptake of cisplatin. Cisplatin also caused decreases of *GST* activity. In rat liver fractions both microsomal and cytosolic *GSTs* were inhibited. As was expected from the glutathione determinations, *GST* was also inhibited in intact rat hepatocytes and in hemolysates made from human erythrocytes and not in intact human eryth-

rocytes. The finding that both human erythrocyte GST (π class) and rat liver cytosolic (mainly α class) and rat liver microsomal GST are inhibited, indicates that the inhibition is rather nonspecific [20] and therefore probably caused by the high direct alkylating activity of cisplatin.

Cyclophosphamide, another well-known alkylating cytostatic drug, was also tested. The alkylating activity of cyclophosphamide is attributed to mustard compounds and acrolein formed as a result of metabolic transformations [13]. In agreement with this need for metabolic activation, cyclophosphamide itself caused only small decreases in the GT content of human red blood cells. This small but significant decrease is interesting because it proves that red blood cells are able to metabolize cyclophosphamide. When rat liver microsomes plus an NADPH regenerating system were added to the red blood cells a strong depletion of intracellular GT was found. This glutathione depletion is accompanied by increased activity of GST in the red blood cells. As expected, cyclophosphamide in the presence of rat liver microsomes and an NADPH regenerating system also caused GT depletion in rat liver cytosol. In this system however, glutathione effectively prevents an effect on GST activity. When diluted cytosol (with lower GSH concentrations) was used an increase in GST activity was found. Activity increases were also found for microsomal GST in the presence of an NADPH regenerating system. This is in good agreement with the findings for the effects on cytosolic GST since no detectable amounts of GT were present in the incubations with microsomes, and GSH protection was therefore excluded. The interpretation of the GST activity increases in human red blood cells and in rat liver cytosol is difficult because of the formation of free radicals by the microsomal system [43, 44]. Addition of a microsomal activating system alone does lead to a decrease in red blood cell and rat liver cytosolic GST activity. Because of this, a possibility exists that the cyclophosphamide dependent activation is really decreased inactivation. Additional experiments showed however that cyclophosphamide addition did not result in lower lipid peroxidation as detected with thiobarbituric acid and did not result in lower methemoglobin formation, and therefore probably does not lead to decreases in free radical release from the microsomes.

Next to cisplatin and cyclophosphamide three other cytostatic drugs were tested. They all belonged to the group of antimetabolites and do not have alkylating activity. Of these 5-fluorouracil and cytosine arabinoside did not affect GT or GST activity in all test systems, while methotrexate only caused an inhibition of microsomal GST. This confirms that the GT depleting activities of cisplatin and of cyclophosphamide metabolites are not a general effect of cytostatic drugs, but can be attributed to the alkylating nature of these substances. The results also suggest that for this class of electrophilic substances the relative effect on GT is larger and more consistent than the effects on GST. The more so because glutathione seems to protect the transferase against low levels of reactive metabolites.

In a third *in vitro* study effects of three industrially used hydroxylamines on

human red blood cells were studied (chapter 5). The parent compound hydroxylamine is known for its hematotoxic potency, with a strong ability to induce methemoglobin [3]. Strong methemoglobin formation was also found for O-ethyl hydroxylamine. Both compounds also induced lipid peroxidation as detectable from the release of thiobarbituric acid reactive compounds. This is probably a result of the formation of free radicals during the oxidation of hemoglobin [34]. As expected the oxidative processes were accompanied by GT depletion in the red blood cells. The relative magnitude of this effect was however much smaller than methemoglobin formation or release of thiobarbituric acid reactive substances. Only a small fraction of the GT that disappeared from the red blood cells could be recovered as oxidized glutathione in the plasma. This indicates that a large fraction of the oxidized glutathione is immobilized within the red blood cells as a result of mixed disulfide formation. This was also found to occur for the hydroxylamine metabolite of the antileprotic drug dapsone [12]. *In vivo* these mixed disulfides will probably be reduced by thioltransferase activity [21] whenever oxidative stress conditions become more favorable. In this way mixed disulfide formation can act as a kind of storage for oxidized glutathione, enabling the cell to retain a nontoxic GSH/GSSG ratio without too much loss of glutathione. While this will be beneficial for the cell it will limit the usability of GT depletion measurements for biological effect monitoring purposes in case of oxidative stress. In parallel with methemoglobin formation and lipid peroxidation, strong decreases in GST activity in red blood cells were found. The same was true for the activity of another protective enzyme, NADPH methemoglobin reductase. Based on these results the use of GST and NADPH methemoglobin reductase measurements for biological effect monitoring of exposure to oxidative agents seems promising.

The third compound used in this *in vitro* study was N,O-dimethyl hydroxylamine. This compound was found to have much less oxidative potency. The methemoglobin formation was much lower than for hydroxylamine or O-ethyl hydroxylamine, there was hardly any lipid peroxidation detectable and we found no GST or NADPH methemoglobin reductase inhibition. On the other hand N,O-dimethyl hydroxylamine addition did result in some GT depletion in the red blood cells, and the increase in plasma GSSG was even somewhat higher than for the other two hydroxylamines. N,O-dimethyl hydroxylamine pretreated red blood cells also had a decreased resistance to hydrogenperoxide induced lipid peroxidation. A likely explanation of this phenomenon was found, when it appeared that N,O-dimethyl hydroxylamine treatment results in a strong inhibition of glucose 6-phosphate dehydrogenase. In the erythrocyte this enzyme is crucial for the production of NADPH. NADPH is needed for many reductive reactions, including glutathione reductase and NADPH methemoglobin reductase activity. Glutathione reductase itself was also, but to a lesser extent, inhibited by N,O-dimethyl hydroxylamine. The reduced availability of NADPH will render the erythrocytes more vulnerable to a subsequent oxidative stress, as was found in the experiments with hydrogenperoxide. This may have practical consequences in

situations where combined exposure to N-substituted hydroxylamines—like N,O-dimethyl hydroxylamine—and compounds with higher direct oxidative potency—like hydroxylamine and O-ethyl hydroxylamine—may occur. Potentiation of the oxidative toxicity by the in itself nonoxidative compounds could be the result of such situations.

Contrary to the conclusions given for electrophilic stress, the hydroxylamine studies indicate that *in vivo* determination of GST (and possibly NADPH methemoglobin reductase) activity in red blood cells may be a more sensitive parameter for oxidative stress than determination of GT depletion. The expected reversibility of mixed disulfide formation *in vivo* will make this even more true.

The first chapter of the second part (chapter 6) gives an overview of a number of biological effect monitoring studies. In a pilot study with smokers and nonsmokers no significant changes in GT concentration or GST activity were found. It became clear from immunochemical determinations however, that the total amount of GST protein was nevertheless increased. Extra synthesis of GST protein must have occurred in smokers without an accompanying increase in GST activity. A possible explanation is that the GST activity is initially decreased as a result of the electrophilic and oxidative stress occurring in smokers. Because of this chronic stress, extra synthesis of GST during erythropoiesis may be induced. This suggests that GST activity in older erythrocytes should be lower than in younger cells. Initially this was contradicted by studies reported in the literature [35], but more recent studies confirm that GST activity is decreased during red cell senescence [11]. This study by Fazi and coworkers is also of interest from another point of view. They proved that erythrocyte GST ρ is converted in another form during red cell senescence. While the K_m value of this secondary form is not changed, it is possible that V_{max} is lowered. The activity loss of GST described in this thesis might involve the same kind of conversion, which according to Shen *et al* [30,31] would be a cysteine modification. This would explain why GST activity seems to stabilize at higher concentrations of oxidative compounds used, as was seen for hydroxylamine and O-ethyl hydroxylamine and, in studies not described here, also for hydrogenperoxide and doxorubicine.

In another study blood and urine samples were collected of two workers preserving wood with coal tar containing creosote oil. Erythrocyte GT concentrations and GST activities in these workers were decreased at the end of the week. After the free weekend, normal values were restored. Despite the very small group size these findings are of interest because of the good correlation with biological monitoring values for coal tar exposure (determined as 1-hydroxypyrene in urine) and genotoxic effects (determined as DNA adducts in white blood cells).

The study described in chapter 7 concerns soil fumigators working in the flower bulb culture in the Dutch "Bollenstreek". This study was carried out in close cooperation with several other institutes. Samples were collected before and after the season during which they applied dichloropropene. At the end of

the season erythrocyte GT concentrations and GST activities had decreased. In clinical chemical studies performed at the universities of Leiden and Amsterdam low levels of liver enzyme induction and subclinical nephrotoxic effects were found in the same group of soil fumigators. The effects on GT and GST were however remarkably less ambiguous than these results from clinical chemical studies. Dichloropropene is a broad working pesticide used for soil fumigation between cultures, and has very high chemical reactivity. As well as the related dichloropropane [14], it is known to be metabolized by glutathione conjugation. In a biological monitoring study, performed at the Free University of Amsterdam, the mercapturic acid derivatives of dichloropropene were detected in urine [40]. The consumption of glutathione by this metabolic activity, is the most likely cause of the decreased erythrocyte GT levels. The decrease in GST activity—which was actually more surprising considering the *in vitro* results described above—must probably be attributed to reactive metabolites and/or reactive breakdown products of cellular constituents.

The biological effect monitoring study described in chapter 8 was directed to what at first sight might be considered a rather special population, which consisted of 23 male and 18 female runners. Previously sedentary, they received intensive running training during a period of 40 weeks. They participated to a competitive run of 15 km length after 20 weeks and ran a half marathon at the end of the study. Blood samples were collected at the start of the study, before and after each run and five days later. After the first 20-week training period GT concentrations and glutathione reductase activity in the erythrocytes were found to have increased. Training consists of periodically high activity with accompanying elevated oxygen turnover. This increased oxygen turnover will result in increased free radical stress [15,17]. The increases in GT and glutathione reductase are probably part of the physiological reaction that counteracts this increased stress. Increases in plasma GSH after prolonged training has also been described in runners [16] and trained beagle dogs [17]. Increased glutathione reductase activities after running training were also reported by Ohno *et al* [25]. The explanation that increased GT values are compensatory for GT loss during acute exercise is consistent with the finding of decreased GT values directly after the 15-km race. After the half marathon the GT values were increased, a phenomenon that is currently hard to explain. Glutathione reductase activities in erythrocytes were increased after both competitive runs. At least a part of these increases was caused by increased availability of the glutathione reductase cofactor riboflavin. Erythrocyte GST activities were lowered after both runs. This is consistent with our expectations, since erythrocyte GST is vulnerable to oxidative stress as it will appear during intensive exercise. As might be expected, the glutathione reductase activities remained elevated after the second training period. In contrast normal GT values were found after 40 weeks. It is not clear whether this is a result of normalized demand for glutathione due to other compensatory mechanisms—like the increased glutathione reductase activity— or whether a lack of synthetical

potential prohibits the preservation of the high GT levels. An indication for the latter explanation was found in the increased GST activities after the second training period. This is probably a reaction to the on going oxidative stress and the associated lipid peroxidation. This suggests that, considering the resulting oxidative effects, the second more intensive training period is less beneficial than the first.

In the final chapter of the experimental part (chapter 9) differences between groups of former Belgian coalminers are described. Both former miners with various stages of coal workers pneumoconiosis ($n = 33$) and former miners without ($n = 58$) this disease were included. It appeared that both erythrocyte GT concentrations and erythrocyte GST activities are decreased during early stages of the inflammatory disease. A causal relationship is likely because in these patients the disease is in an active phase. During continuous inflammatory activity large portions of active oxygen species are generated by the alveolar macrophages [28]. These active oxygen species are expected to be responsible for the effects on GT and GST found. Also in the sporting population a more diffusive form of inflammatory activity may have contributed to the oxidative effects seen.

It can be concluded that the two most important parameters described in this thesis —the glutathione concentration and the glutathione S-transferase activity in red blood cells— are interesting tools for mechanistic toxicological research and for biological effect monitoring. In in vitro experiments glutathione seems to give the most interesting results in situations where electrophilic exposure is at stake. In vivo effects were also seen after heavy exercise and in miners, and were reported in the literature as a result of lipid peroxidation for workers exposed to lead [36]. Glutathione S-transferase activity is of most interest in situations where oxidative stress occurs. It may be used as an interesting addition to, and an alternative for, the problematic determination [27] of in vivo lipid peroxidation.

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Hoofdstuk 11

Samenvatting en Discussie

Chemische factoren die een risicobron vormen voor beschadigingen van belangrijke biologische macromoleculen, waaronder het genetisch materiaal, laten zich ruwweg in twee groepen verdelen. Enerzijds zijn er de electrofiele verbindingen, en stoffen die tot dergelijke verbindingen kunnen worden gemetaboliseerd. Deze stoffen zijn direct in staat biologische macromoleculen te beschadigen. Daarbij treedt dan veelal adductvorming op. Anderzijds zijn er omstandigheden die leiden tot de vorming van vrije radicalen. Dergelijke vrije radicalen kunnen eveneens tot beschadiging van biologische materialen leiden, zij kunnen echter ook aanleiding geven tot aantasting van de lipide structuur van membranen via lipide-peroxidatie. Bij deze lipide-peroxidatie ontstaan opnieuw reactieve verbindingen, die op hun beurt weer elders schade kunnen aanrichten. Het in dit proefschrift besproken glutathion – glutathion S-transferase systeem vormt een bijzondere factor doordat het zowel een bijdrage levert aan de bescherming tegen electrofiele als aan het verminderen van oxidatieve stress.

Bescherming tegen electrofiele verbindingen komt tot stand door koppeling van de nucleofiele thiolgroep van het glutathion aan het electrofiele reagens. In veel gevallen verloopt deze reactie spontaan, zij kan echter ook door het glutathion S-transferase worden gekatalyseerd. De zo gevormde thioether-produkten worden in het algemeen uit de cel afgevoerd, en verder gemetaboliseerd. In principe kan dit leiden tot een vermindering van het in de cel aanwezige glutathion. Bovendien is reeds langere tijd bekend dat in sommige gevallen reactieve verbindingen niet alleen via de S-transferase reactie worden gekoppeld, maar ook via een soort zelfmoordreactie aanleiding geven tot een beschadiging van het glutathion-S-transferase zelf.

De bescherming tegen oxidatieve stress is complexer van aard. Glutathion kan bijdragen aan de enzymatische reductie van vitamine E, zowel direct als indirect via de reductie van vitamine C. Het goed vetoplosbare, gereduceerde vitamine E beschermt effectief tegen lipide-peroxidatie processen. Als gevolg van oxidatieve stress treedt ook oxidatie van eiwit-sulhydrylgroepen op. Dit kan leiden tot crosslinking van eiwitten. Hierdoor kunnen enerzijds enzymactiviteiten verloren gaan, en kunnen anderzijds, door crosslinking van structuureiwitten,

vormveranderingen van cellen zonder cytoskelet optreden. Van rode bloedcellen, die voor hun mobiliteit door perifere weefsels in sterke mate afhankelijk zijn van de vervormbaarheid van hun celmembraan, is bekend dat zij vatbaar zijn voor dit soort oxidatie processen. Glutathion kan via thioltransferase activiteit gebruikt worden om geoxideerde sulfhydrylgroepen te reduceren. Ook kan als gevolg van de oxidatieve stress crosslinking van de glutathion-sulfhydrylgroep met eiwit-sulfhydrylgroepen optreden. Met name in rode bloedcellen is gereduceerd glutathion ook een algemene buffer voor reducerend vermogen. In deze cellen zijn namelijk geen mitochondriën aanwezig, die in andere celtypes voor de snelle levering van grotere hoeveelheden energie en reductie-equivalenten kunnen zorgen. Homeostase van de glutathion redox status wordt, bij voldoende aanlevering van glucose, gegarandeerd via regulatie van de glutathion reductase en de glucose 6-fosfaat dehydrogenase activiteit.

Bovendien is glutathion van belang in de glutathion-peroxidase reactie. Gekatalyseerd door het selenium afhankelijke enzym glutathion-peroxidase ontgift het zowel waterstofperoxide als de bij lipide-peroxidatie gevormde organische peroxiden. De ontgifting van de laatste categorie peroxiden kan ook worden gekatalyseerd door glutathion S-transferases. Deze activiteit van glutathion S-transferase wordt ook wel aangeduid met selenium onafhankelijke glutathion-peroxidase activiteit. De genoemde beschermende reacties leiden in alle gevallen tot de oxidatie van glutathion. In principe kan dit geoxideerde glutathion vervolgens door glutathion-reductase weer worden gereduceerd. Geoxideerd glutathion is echter toxisch voor rode bloedcellen, en wordt hieruit actief verwijderd. De oxidatieve vorming van glutathion-eiwitbruggen, ook wel aangeduid met mixed disulfides, leidt bovendien direct tot een verlies van beschikbaar glutathion. Dit betekent dat, net als bij electrofiële stress, ook als gevolg van oxidatieve stress een afname van de beschikbare hoeveelheid glutathion kan optreden. De analogie in effecten van beide stress-factoren gaat echter nog verder. Het is bekend dat het glutathion S-transferase iso-enzym dat aanwezig is in de rode bloedcel, en dat behoort tot de zogenaamde π klasse, zelf gedeeltelijk geïnactiveerd kan worden als gevolg van oxidatieve stress.

Het bovenstaande, grotendeels gebaseerd op in de literatuur beschreven in vitro experimenten, geeft aan dat het a priori denkbaar was dat als gevolg van een belasting met electrofiële of oxidatieve xenobiotica of als gevolg van een op andere wijze tot stand gekomen oxidatieve stress, veranderingen in het glutathion-gehalte of in de glutathion S-transferase activiteit van rode bloedcellen zouden optreden. De studies beschreven in dit proefschrift waren er op gericht de praktische bruikbaarheid van dit idee voor experimenteel toxicologisch onderzoek en voor biologisch effect monitoring te onderzoeken. Het eerste deel (hoofdstuk 3-5) bevat in vitro studies met humaan bloed. In het tweede deel (hoofdstuk 6-9) wordt een aantal in vivo biologisch effect monitoring studies beschreven.

Direct reactieve electrofiële verbindingen zoals iodoaceetamide, diethylma-

leaat en N-ethylmaleimide zullen, ongeacht de aanwezigheid van al dan niet intacte bloedcellen, in sterke mate met glutathion reageren. De aanwezigheid van glutathion in de rode bloedcel werkt daardoor in feite beschermend t.o.v. andere biologische effecten. Uit hoofdstuk 3 blijkt dat voor deze zeer reactieve verbindingen de vorming van hemoglobine-adducten sterk toeneemt nadat het glutathion volledig is verbruikt. Beïnvloeding van de effectiviteit van de glutathion bescherming *in vivo* zal ook consequenties hebben voor de mate van hemoglobine-adduct vorming. Dit is mogelijk van belang omdat hemoglobine-adduct nivo's zelf soms gehanteerd worden als biologisch effect parameter voor expositie aan electrofiele genotoxische agentia.

Uit hoofdstuk 4 blijkt dat ook het direct alkylerende cytostaticum cisplatina *in vitro* depletie van glutathion veroorzaakt in geïsoleerd cytosol van humane rode bloedcellen en in al dan niet intacte rattelever-cellen. In intacte rode bloedcellen vertoont cisplatina dit effect niet, waarschijnlijk doordat het celmembraan van dit type cellen opname effectief voorkomt. Cisplatina veroorzaakte naast glutathion-depletie ook een inhibitie van glutathion S-transferase. Deze inhibitie werd na belasting van rattelever-fracties zowel gevonden voor de enzymactiviteit in microsomen als voor die in cytosol. In overeenstemming met wat werd gevonden voor de glutathion-gehaltenes veroorzaakte cisplatina in intacte humane rode bloedcellen geen effect op glutathion S-transferase en werd wel een inhibitie gevonden wanneer cisplatina aan hemolysaten of aan al dan niet intacte hepatocyten werd toegevoegd. De inhibitie van glutathion S-transferase door cisplatina blijkt dus niet iso-enzym specifiek te zijn, en is waarschijnlijk een direct gevolg van de hoge alkylerende activiteit van dit cytostaticum.

Een tweede alkylierend cytostaticum, cyclophosphamide, werd eveneens getest. Van cyclophosphamide is bekend dat het moet worden gemetaboliseerd alvorens het zijn biologische activiteit kan uitoefenen. In overeenstemming hiermee bleek cyclophosphamide zelf slechts geringe effecten te hebben op het glutathion-gehalte van rode bloedcellen. Dit geringe, maar significante effect zelf is interessant omdat het aangeeft dat cyclophosphamide metabolisme ook kan optreden in de rode bloedcel. In aanwezigheid van metabole systemen op basis van rattelever-microsomen geeft cyclophosphamide een sterke depletie van het glutathion in rode bloedcellen. Deze depletie gaat gepaard met een activatie van het glutathion S-transferase in deze cellen. In aanwezigheid van een metabool systeem op basis van rattelever-microsomen veroorzaakt cyclophosphamide, zoals verwacht, ook een depletie van het glutathion in rattelever-cytosol. Hier voorkomt glutathion echter effectief een effect op het glutathion S-transferase: slechts bij gebruik van verdund cytosol werd een verhoging van deze enzym activiteit gevonden. De activiteit van het microsomale glutathion S-transferase werd, mits een NADPH regenererend systeem aanwezig was, wel verhoogd door cyclophosphamide. In dit testsysteem waren echter geen aantoonbare hoeveelheden glutathion aanwezig, zodat een bescherming door glutathion was uitgesloten. De interpretatie van de glutathion S-transferase activeringen in rode bloedcellen en rattelever-cytosol

wordt bemoeilijkt doordat het microsomale systeem zelf, doordat vrije radicalen gevormd worden, een inactiverend effect op de glutathion S-transferase activiteit heeft. Het is mogelijk dat er feitelijk geen activatie, maar een verminderde inactivatie plaats vindt. Aanvullende experimenten gaven daarbij echter wel aan dat dit dan niet gepaard gaat met een verminderde radicaal vorming zoals die valt te detecteren in de vorm van verbindingen die reactief zijn t.o.v. thiobarbituurzuur of die aanleiding geven tot methemoglobine vorming.

Drie andere cytostatica, allen zogenaamde anti-metaboliëten, werden eveneens getest. 5-Fluoruracil en cytosine-arabinoside hadden geen enkel effect op de glutathion-gehalten en de glutathion S-transferase activiteiten in de diverse geteste systemen. Methotrexate veroorzaakte uitsluitend een inhibitie van het microsomale glutathion S-transferase. Dit bevestigt dat de voor cisplatina en cyclophosphamide metaboliëten gevonden glutathion depletende eigenschappen niet een algemeen effect van cytostatica zijn, maar eerder een gevolg zijn van het alkylerend karakter van deze verbindingen. Ook suggereren de resultaten van de experimenten met cyclophosphamide en cisplatina dat voor electrofiële verbindingen het relatieve effect op glutathion groter, en meer eenduidig is dan dat op glutathion S-transferase. Dit te meer omdat glutathion zelf ook het transferase lijkt te beschermen tegen lage concentraties aan electrofielen zoals die bijvoorbeeld via de metabole activatie van cyclophosphamide ontstaan.

In een derde *in vitro* studie werden de effecten van een drietal industriële chemicaliën die behoren tot de zogenaamde hydroxylamines (HO-N-derivaten) bestudeerd. Van de moederverbinding, hydroxylamine, was reeds bekend dat zij hematotoxisch is en ondermeer methemoglobine veroorzaakt. Sterke methemoglobine-vormende eigenschappen werden ook gevonden voor het O-ethylhydroxylamine. Beide verbindingen gaven, waarschijnlijk als gevolg van de vrije radicalen die ontstaan bij de methemoglobine-vorming, bovendien lipide-peroxidatie. Zoals verwacht werd onder deze oxidatieve omstandigheden ook een vermindering van het beschikbare glutathion in de rode bloedcellen gevonden. Dit effect was echter beduidend minder sterk dan de methemoglobine-vorming en de lipide-peroxidatie zelf. Opvallend was dat het verdwenen glutathion slecht voor een klein deel in geoxideerde vorm in het plasma werd teruggevonden. Kennelijk is veel glutathion gebonden aan eiwitten. *In vivo* kunnen dergelijke mixed disulfides na vermindering van de oxidatieve stress weer gereduceerd worden, hetgeen de bruikbaarheid van glutathion-depletie voor biologisch effect monitoring zou kunnen beperken. Parallel aan de methemoglobine-vorming en lipide-peroxidatie werd een sterke vermindering van de glutathion S-transferase activiteit in de rode bloedcellen gevonden. Hetzelfde gold overigens voor de activiteit van een ander enzym, het NADPH-methemoglobine-reductase. Op basis van deze bevindingen lijkt het interessant deze beide parameters te testen in biologisch effect monitoring studies met oxidatieve verbindingen. Een derde hydroxylamine verbinding, het N,O-dimethylhydroxylamine, bleek aanzienlijk minder oxidatieve capaciteit te hebben. Er werd aanzienlijk minder methemoglobine-vorming, nau-

welijks lipide-peroxidatie, en in het geheel geen glutathion S-transferase remming gevonden. Wel bleek ook deze verbinding aanleiding te geven tot milde glutathion-depletie in rode bloedcellen, waarbij zelfs een wat sterkere toename van de hoeveelheid glutathion in plasma werd gevonden. Bovendien bleken N,O-dimethylhydroxylamine behandelde bloedcellen een duidelijk verminderde weerstand tegen door waterstofperoxide geïnduceerde lipide-peroxidatie te hebben. Een waarschijnlijke verklaring voor deze fenomenen werd gevonden in de inhibitie van glucose-6-fosfaat-dehydrogenase door N,O-dimethylhydroxylamine. Verlies van deze enzymactiviteit maakt dat de cel niet meer kan beschikken over NADPH dat nodig is voor reducerende reacties. Daardoor zou ondermeer het glutathion reductase niet meer kunnen functioneren. Het glutathion reductase werd daarnaast ook zelf geremd door N,O-dimethylhydroxylamine. Maar deze inhibitie was minder sterk. De verminderde weerstand tegen oxidatieve stress, zoals veroorzaakt door het zelf minder sterk oxidatieve N,O-dimethylhydroxylamine, zou van praktische betekenis kunnen zijn omdat bij gecombineerde blootstelling aan N,O-dimethylhydroxylamine en bijvoorbeeld O-ethylhydroxylamine potentiëring van de toxiciteit van die laatste het gevolg zou kunnen zijn.

In tegenstelling tot de hierboven beschreven conclusies voor electrofiele verbindingen geven de hydroxylamine-studies aan dat bij oxidatieve belasting de activiteit van glutathion S-transferase (en mogelijk ook die van het NADPH-met-hemoglobine-reductase) een gevoeliger parameter is dan de glutathion-depletie. Dit is extra interessant omdat glutathion-depletie via oxidatie voor een belangrijk deel blijkt te bestaan uit de vorming van eiwit-glutathion disulfiden. Deze gemengde disulfiden zullen bij voldoende beschikbaarheid van NADPH-reductie-equivalenten opnieuw gereduceerd worden via een gecombineerde activiteit van thioltransferase en glutathion-reductase. Het valt te verwachten dat een dergelijk compensatoir mechanisme de relatieve betekenis van glutathion oxidatie *in vivo* nog zal verminderen.

In het eerste hoofdstuk van het tweede deel (hoofdstuk 6) wordt een overzicht gegeven van een aantal biologisch effect monitoring studies. In een pilot studie met rokers en niet-rokers werd geen aantoonbaar effect van roken op de glutathion S-transferase activiteit gevonden. Uit immunochemische bepalingen kwam echter naar voren dat de totale hoeveel glutathion S-transferase eiwit wel was verhoogd. Kennelijk is er bij de rokers extra glutathion S-transferase gesynthetiseerd, zonder dat de enzymactiviteit is toegenomen. Een mogelijke verklaring is dat als een gevolg van de zowel oxidatieve als electrofiele stress die door het roken wordt veroorzaakt in eerste instantie de enzymactiviteit afneemt, waarbij vanwege de chronische belasting ter compensatie tijdens de erythropoiese extra glutathion S-transferase wordt aangemaakt. Een tweede studie had betrekking op een tweetal werknemers betrokken bij de houtconservering m.b.v. koolteer houdende creosoot-olie. Bij deze werknemers werden aan het eind van de werkweek verlaagde glutathion-gehalten en verlaagde glutathion S-transferases activiteiten

in rode bloedcellen aangetoond. Na het weekend waren deze waarden weer hersteld. Ondanks de zeer kleine groepsomvang is deze bevinding interessant omdat gevonden effecten op de beide hier bestudeerde parameters goed bleken te correleren met biologische monitoring waarden voor koolteerbelasting (gemeten als hydroxypyreen in urine) en met genotoxische effecten (vastgesteld via bepaling van DNA-adducten in witte bloedcellen).

Hoofdstuk 7 beschrijft een studie bij een veertiental grondontsmetters afkomstig uit de Bollenstreek, die tijdens de onderzoeksperiode werkten met dichloorpropeen. Deze studie werd gezamenlijk met diverse andere instituten uitgevoerd. Na afloop van het seizoen bleek zowel de glutathion als de glutathion S-transferase waarde bij de twaalf op deze parameters onderzochte grondontsmetters duidelijk verlaagd te zijn. In aan de universiteiten van Leiden en Amsterdam uitgevoerde klinisch chemische studies werden indicaties gevonden voor een geringe leverenzym-inductie en voor een subklinisch toxisch effect op de nier. Opvallend was dat de effecten op glutathion en glutathion S-transferase aanzienlijk eenduidiger waren dan de resultaten van deze klinisch chemische studies. Dichloorpropeen is een zogenaamde allesdoder met een zeer hoge reactiviteit. Van deze stof bekend is dat zij gedeeltelijk via de mercaptuurzuur-route wordt gemetaboliseerd. In aan de Vrije Universiteit uitgevoerde biologische monitoring studies bij deze zelfde werknemers, werden in urine ook inderdaad verhoogde gehalten aan mercaptuurzuur-metabolieten van dichloorpropeen aangetoond. Het glutathion-verbruik tijdens dit metabolisme is waarschijnlijk primair verantwoordelijk voor de optredende glutathion-depletie. De afname van de glutathion S-transferase activiteit valt waarschijnlijk toe te schrijven aan de gevormde reactieve metabolieten en/of cel-afbraakprodukten.

In hoofdstuk 8 wordt een biologisch effect monitoring studie bij een op het eerste gezicht wat bijzondere populatie beschreven. Het betreft hier geen beroepsmatig blootgestelden, maar 23 mannelijke en 18 vrouwelijke sporters. Ongetrainde individuen werden gedurende een langere periode intensief getraind. Na 20 weken namen zij deel aan een 15 km hardloop wedstrijd, en na 40 weken aan een halve marathon. Aan het begin van de studie, voor en na de beide wedstrijden, en ook enkele dagen daarna werden bloedmonsters afgenomen. Tijdens de eerste 20 weken trainingsperiode nam zowel het glutathion-gehalte als de glutathion-reductase activiteit in de rode bloedcellen toe. Mogelijk zijn deze beide toenames het gevolg van een behoefte aan extra bescherming tegen de hogere zuurstof-turnover en daarmee gepaard gaande oxidatieve stress. In overeenstemming hiermee werd direct na de 15 km wedstrijd een tendens richting lagere glutathion-waarden waargenomen. Na de halve marathon was het glutathion-gehalte overigens, om niet geheel duidelijke redenen, verhoogd. Het glutathion-reductase was direct na de beide wedstrijden actiever dan daarvoor, iets wat op zijn minst gedeeltelijk werd veroorzaakt door de beschikbaarheid van extra riboflavine. Glutathion S-transferase was na beide wedstrijden in activiteit verlaagd. Dit is in overeenstemming met de verwachting. Immers het transferase is gevoelig voor de extra oxidatieve stress,

zoals die tijdens inspanning zal optreden. Na de tweede trainingsperiode was het glutathion-reductase volgens verwachting nog altijd verhoogd. Dit gold echter niet voor het glutathion. Het is daarbij op zichzelf niet duidelijk of dit een gevolg is van verminderde behoefte als gevolg van andere aanpassingen (zoals de toegenomen reductase activiteit) of van gebrek aan synthese capaciteit. Een indicatie in deze laatste richting wordt wel gegeven door de bevinding dat de glutathion S-transferase activiteit na deze tweede trainingsperiode was gestegen. Waarschijnlijk is dit een reactie op de voortdurende oxidatieve stress, en de daarmee waarschijnlijk gepaard gaande lipide-peroxidatie. Dit wijst er op dat vanuit het oogpunt van oxidatieve belasting de tweede, intensievere trainingsfase minder positief is dan de eerste.

In hoofdstuk 9 tenslotte worden verschillen beschreven tussen diverse groepen Belgische ex-mijnwerkers met ($n = 33$) en zonder ($n = 58$) mijnwerkers pneumoconiose. Het blijkt dat tijdens de vroege stadia van de inflammatoire ziekte zowel het glutathion als het glutathion S-transferase verlaagd zijn. Een oorzakelijk verband is waarschijnlijk omdat het hier gaat om mensen met actieve stadia van deze ziekte. Tijdens de voortdurende inflammatoire activiteit worden door de alveolaire macrofagen grote hoeveelheden reactieve zuurstofspecies gevormd. Het is waarschijnlijk dat die uiteindelijk verantwoordelijk zijn voor de beschreven effecten. Ook bij de sporters kan een als gevolg van spierbeschadigingen optredende meer diffuse vorm van inflammatie een rol hebben gespeeld.

Geconcludeerd kan worden dat de beide belangrijkste beschreven parameters, het glutathion en de glutathion S-transferase activiteit in rode bloedcellen, interessante gereedschappen vormen voor mechanistisch onderzoek en voor biologisch effect monitoring. Glutathion lijkt daarbij het meest relevant in die gevallen waarin het gaat om electrofiele belastingen. Glutathion S-transferase daarentegen is vooral van betekenis wanneer het gaat om oxidatieve belastingen. Het kan daarbij een bruikbare aanvulling vormen op de, op zichzelf nog altijd problematische, detectie van in vivo lipide-peroxidatie.

Curriculum Vitae

Ik werd op 7 april 1957 geboren te Gouda. Een stadje waar ik me vooral van herinner dat de grond er altijd langzaam onder vandaan leek te zakken. Omdat de eerste jaren vaders wil nu eenmaal wet was, bezocht ik de lagere school in diverse kleine West-Duitse dorpjes. In 1969 begon ik mijn middelbare school loopbaan op het internaat van het Gabriel College in Mook. Twee jaar later bleek dit een onhoudbare situatie en werd het internaat opgeheven. Het diploma Atheneum B behaalde ik in 1975 dan ook aan Christelijk Lyceum — met een harde G — te Alphen aan den Rijn. Ik studeerde vervolgens Moleculaire Wetenschappen aan de Landbouw Hogeschool te Wageningen, waar de Wageningse Lente toevallig tijdens mijn aanwezigheid bleek te ontluiken. Na wat bestuurlijke activiteiten en een jaar als werkstudent projektonderwijs studeerde ik in 1983 af. In die tijd was het nog mogelijk een vakkenpakket te kiezen. Het werden hoofdvakken Biochemie (bij Dr. H. Haaker) en Toxicologie (bij Dr. B. Blaauboer in Utrecht), en een praktijkperiode bij de Vereniging Milieudefensie. De activiteiten daar hadden, anders dan de titel "Sputten kan niet meer" doet vermoeden, geen betrekking op hard drugs maar op pesticiden gebruik in openbaar groen. In het bijvak wetenschapsfilosofie tenslotte vond ik de ruimte ook in studietijd over de zin en onzin van dat alles na te denken.

Na te zijn afgestudeerd kwam ik in dienst bij wat later de vakgroep Toxicologie van de Katholieke Universiteit Nijmegen zou worden, eerst als wetenschappelijk assistent en later als universitair docent. Het daar uitgevoerde onderzoek had betrekking op de vorming van hemoglobine adducten na beroepsmatige blootstelling aan genotoxische verbindingen. Het was er op gericht eenvoudige ook in de bedrijfsgezondheidszorg bruikbare methodieken te ontwikkelen. Hoewel er natuurlijk nog ongekende mogelijkheden moeten liggen in het gebruik van een stethoscoop, bleek dit uitgangspunt helaas een misvatting. Gelukkig kon, door het bestuderen van de invloed die glutathion heeft op de hemoglobine adduct vorming, in die tijd ook al een aanzet worden gegeven voor het in dit proefschrift beschreven onderzoek. In 1989 kwam ik in dienst bij de vakgroep Arbeidsgeneeskunde, Milieugezondheidskunde en Toxicologie van de Rijksuniversiteit Limburg. In een ten gevolge van reorganisaties roerige periode, als uitvloeisel waarvan de vakgroep uiteindelijk hetzelfde lot was beschoren als het Gabriel College, werden ondermeer de veldstudies met marathon lopers en mijnwerkers zoals beschreven in dit proefschrift afgerond. Vanaf eind 1991 kwam ik als leider van de Toxicologie sectie bij de vakgroep Farmacologie in een wat rustiger vaarwater. Hier werd nog een deel van de beschreven *in vitro* studies uitgevoerd, en vond uiteindelijk de afronding van dit proefschrift plaats.

Chris Evelo.

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Nawoord

Een heleboel mensen zijn op de een of andere manier betrokken geweest bij het tot stand komen van dit proefschrift. Een aantal daarvan wil ik hier expliciet bedanken, gewoon omdat ik dat belangrijk vindt. Ongetwijfeld zal ik er ook een paar vergeten, niet omdat zij niet belangrijk zijn geweest, maar omdat ik er nu even niet aan denk. Ik hoop dat zij me dat vergeven.

Mijn promotor prof. Henderson, die mij toch maar aan twee verschillende universiteiten een aanstelling bezorgde, voor het vertrouwen, en voor de vele gesprekken, juist omdat die lang niet allemaal over wetenschap gingen.

Alle collega's van de vakgroep Toxicologie in Nijmegen — met name ook die van wat voor mij het eerste uur was — voor de prettige sfeer. Hennie Roelofs, Janny Peters en Peter Fijneman zijn daarbij van onschatbaar belang geweest door het vele praktische werk. Rob Bos, Frans Jongeneelen, en John Neis lieten mij drie heel verschillende manieren van onderzoek doen zien, ik heb daar veel van geleerd. Frans was bovendien de eerste die filosofeerde over de mogelijkheid om biomonitoring van thioethers te combineren met onderzoek naar effecten op het systeem waarin die thioethers worden gevormd. Mogelijk een cruciale stap op weg naar het hier beschreven onderzoek.

De moeilijke tijd die ik meemaakte bij de voormalige vakgroep Arbeidsgeneeskunde, Milieugezondheidkunde en Toxicologie had nooit zo gezellig kunnen zijn zonder het relativerend vermogen van mijn collega's, bedankt daarvoor. Wat overigens te denken van de opmerking: "och, voordat jij kwam was het eigenlijk allemaal nog niet zo erg ...".

De vakgroep farmacologie, dat zijn er meteen een heleboel, maar het is wel gewoon een goeie club. En je moet het toch maar willen, zo'n stelletje dakloze toxicologen opnemen. Daarvoor ben ik met name prof. Struijker Boudier, mijn huidige vakgroep voorzitter, bijzonder dankbaar. De helderheid waarmee afspraken worden gemaakt vind ik nog steeds een verademing. Vakgroepen, zo groot als farmacologie zitten niet zomaar gevangen in het web van de universitaire structuur, zij spinnen ook zelf behoorlijk mee. Dat was voor mij een nieuwe ervaring, en een hele leerzame.

De mensen van de sectie Toxicologie, Nicole Palmen, Leo Baars, Anita Spooren en alle studenten die daar de afgelopen paar jaar een poosje bij wilden horen, kortom mijn eigen groep. Jullie waren in die periode eigenlijk het allerbelangrijkst. Door jullie was het fijne tijd. Jullie zullen ook wel het blijst zijn dat dit proefschrift er nu is, en er weer andere dingen centraal komen te staan.

Heel veel studenten hebben stukjes onderzoek uitgevoerd die voor mij op dat moment een deel van de legpuzzel vormden. Lianne de Wijs, Geert-Jan Reimer, Toon van Bommel, Karel Herberichs, Anita Spooren en Rob Bisschops zullen hun eigen puzzelstukjes in dit proefschrift hebben terug gezien. Een heleboel andere stukjes pasten uiteindelijk niet zo goed. Hele onderzoekslijnen, bijvoorbeeld

op de gebieden hemoglobine adducten, biomonitoring van cytostatica en fysiologische simulatie modellen, zijn zelfs buiten beeld gebleven.

The discussions with Dr. Niemeyer and Dr. Zechel from Asta about the special aspects of cyclophosphamide proved to be of importance, not only for the biomonitoring studies described elsewhere, but also for the effect studies described in this thesis. Marjo Catsberg van het Maasland Ziekenhuis deed de Heinz Body analyse in de hydroxylamine studie, een erg ondankbare taak. John Neis en Jan Bakker van DSM waren niet alleen als opdrachtgever betrokken bij deze studie. Hun kritische houding, praktijk gerichtheid en de behoefte om echt mee te denken maakten de samenwerking tot een genoegen. Van Wilbert Peters, Ria Vos en prof. van Bladeren heb ik in de korte tijd dat we een soort regionaal GST overleg hadden een heleboel zinnige tips gekregen; toch jammer dat Maastricht niet zo in die regio ligt.

Voor al bij de monitoringsstudies beschreven in het tweede deel van dit proefschrift zijn een heleboel mensen betrokken geweest. Veel van het organisatorische werk in de pilot studie met creosoteerders was in handen van Frans Jongeneelen. Rob Baan deed de bijbehorende DNA-adduct metingen. De organisatie van de dichloorpropeen studie was in handen van Ellie Brouwer, die samen met Ton Verplanke bovendien de lever- en nierfunctie analyses uitvoerde. Ronald van Welie deed de specifieke mercaptuurzuur analyses. De professoren de Wolff en Vermeulen hadden bovendien een belangrijke inhoudelijke inbreng in deze studies. Beiden zijn zij ook op diverse andere momenten door hun grote interesse en creativiteit een belangrijke stimulans geweest. Gène Janssen zorgde op zijn eigen unieke wijze voor de organisatie van de marathon studie. Zonder hem zou deze studie niet uitvoerbaar zijn geweest. Dr. Artur of the Center for Preventive Medicine in Nancy was involved in the study of the oxidative aspects of physical activity. I hope to continue this collaboration in the near future with respect to the antioxidant enzymes. Voor de mijnwerkers studie was al het organisatorische werk al verzorgd door Paul Borm. In diverse gesprekken over allerlei aspecten van het onderzoek bleek Paul telkens weer verrassende ideeën te hebben. Dat hielp zeker om er ook zelf steeds weer vanuit een andere hoek tegenaan te kijken.

Zonder de medewerking van de staf en medewerkers van de Rode Kruis bloedbanken in Nijmegen en Maastricht was dit proefschrift waarschijnlijk ten onder gegaan aan bloedarmoede.

Tenslotte wil ik de medewerkers van de UB in Maastricht bedanken voor hun extreem goede service. Boeken uit het magazijn zijn er altijd sneller dan beloofd (dat kan natuurlijk ook een slimmigheidje zijn), en soms slagen ze er in publicaties te vinden die ik op basis van mijn zelf ingevulde informatie nooit boven tafel zou hebben gekregen.

Chantal, Jannemiek en Sjoerd herinnerden mij er, ieder op een eigen manier, telkens weer aan dat er nog zoveel meer is.

TABLE II. Results of the LRT method in computer simulations of markers attached to a moving left ventricle.

$\sigma^{1)}$ pixel	$d_m^{2)}$ pixel	$M=25^{3)}$		$M=50$		$M=75$		$M=100$	
		$f_{DE}^{4)}$	$f_{CL}^{5)}$	f_{DE}	f_{CL}	f_{DE}	f_{CL}	f_{DE}	f_{CL}
0.5	4	4	99	7	95	9	91	14	88
0.5	8	10	97	25	90	38	84	52	79
1.0	4	3	92	7	84	9	78	15	71
1.0	8	10	90	25	80	40	69	-	-

1) RMS noise level of the marker image position measurement

2) Minimum distance allowed between marker image positions

3) Number of markers used in the simulations

4) The percentage of the $M \times 60$ in 60 frames generated marker image positions, which was deleted because positions were mutually closer than d_m

5) The percentage of the retained marker image positions which belonged to correct reconstructed tracks of a duration of at least 50 frames

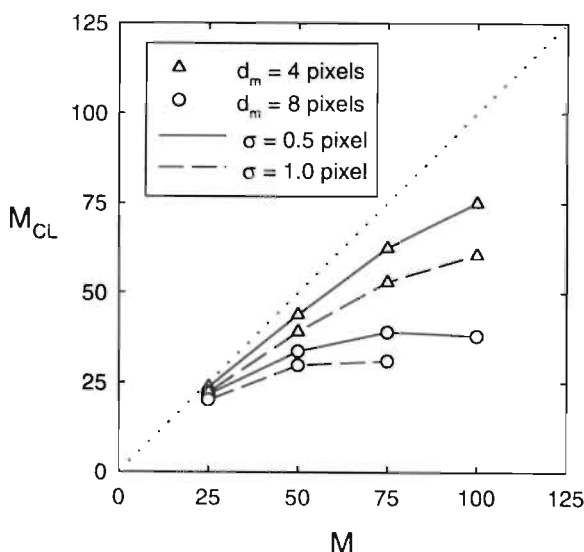


FIGURE 6. The performance index M_{CL} is the average number of marker image positions per frame belonging to a correct track with a duration of at least 50 frames. M_{CL} is shown as a function of the number of markers M . The computer simulations were carried out with a minimum distance allowed between marker image positions, d_m , of 4 (triangle) and 8 pixels (circle), and the RMS noise level σ equal to 0.5 (solid line) and 1 pixel (dashed line), respectively. The dotted line $M_{CL}=M$ refers to perfect performance.